



Present and future of gene therapy in Neuromuscular Diseases

Satellite Scientific Symposium endorsed by ERN EURO-NMD

February, 22nd 2024

Working Group
Neuropathies

Working Group
Gene Therapies

Gene Therapies for peripheral neuropathies *Present and future*

Kleopas A. KLEOPA, MD, PhD, FAAN, FEAN

NEUROMUSCULAR DISORDERS CENTER

& Department of Neuroscience

THE CYPRUS INSTITUTE OF NEUROLOGY AND GENETICS

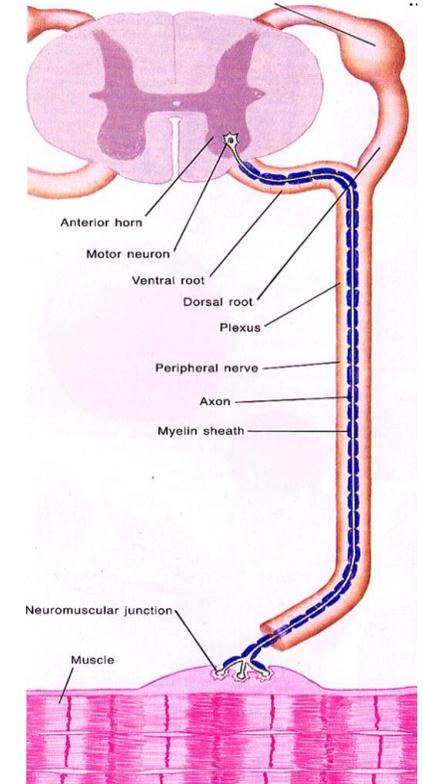
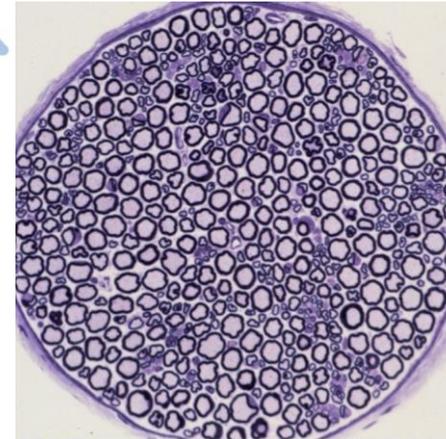
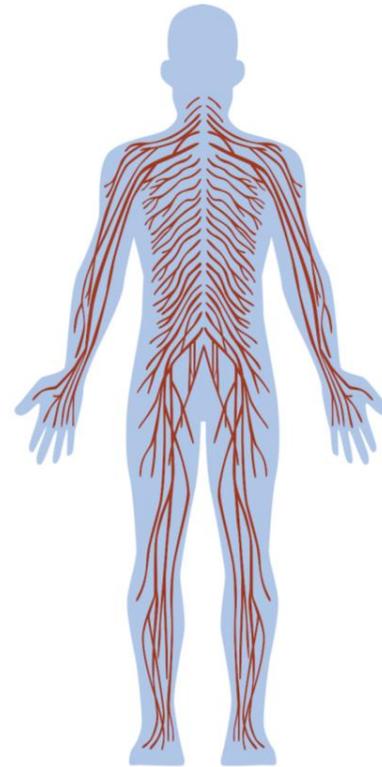
Nicosia, CYPRUS



Gene therapy developments in peripheral neuropathies

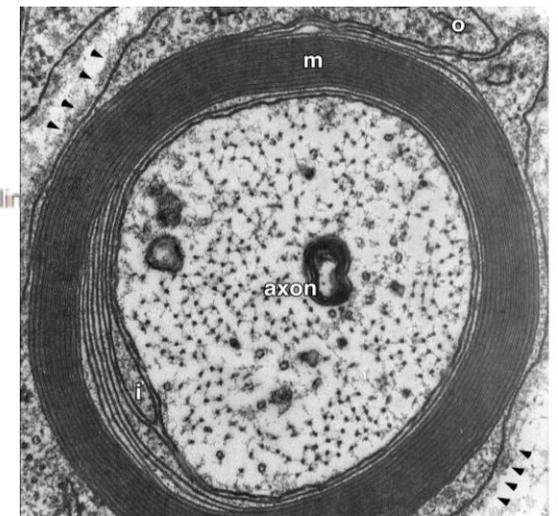
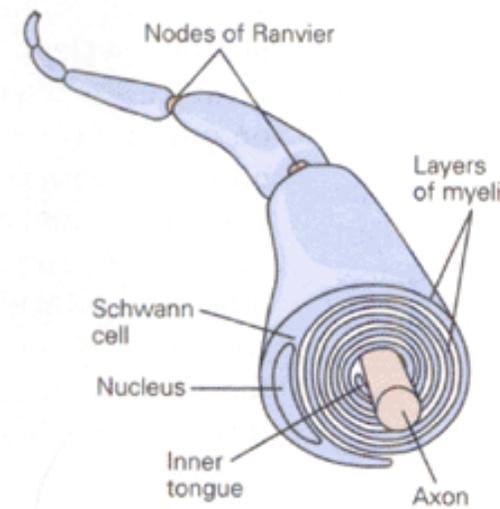
Syndromic neuropathies

- Hereditary TTR variant amyloidosis (hATTRv)
- Giant axonal Neuropathy (GAN)



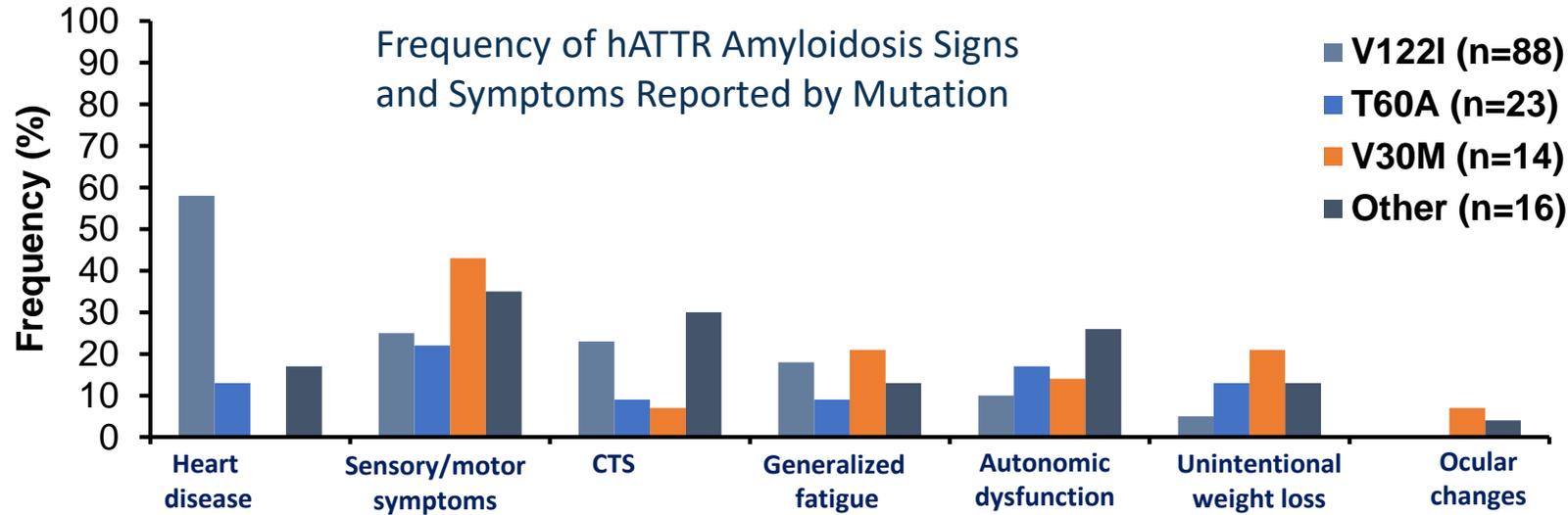
Non-syndromic neuropathies

- Charcot-Marie-Tooth (CMT) diseases
 - Demyelinating CMT Types 1 and 4
 - Axonal CMT Type 2
 - Intermediate CMT

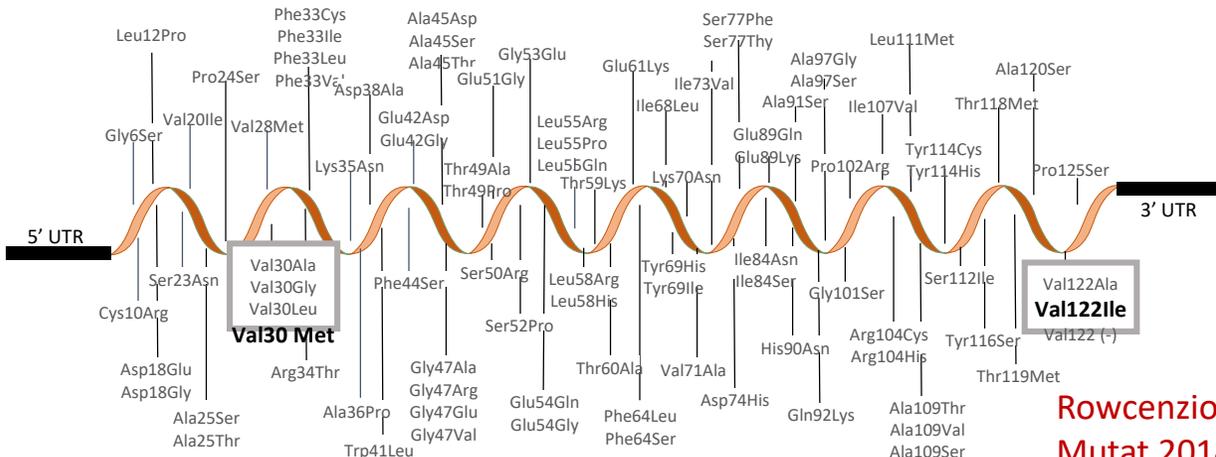


hATTRv Amyloidotic polyneuropathy

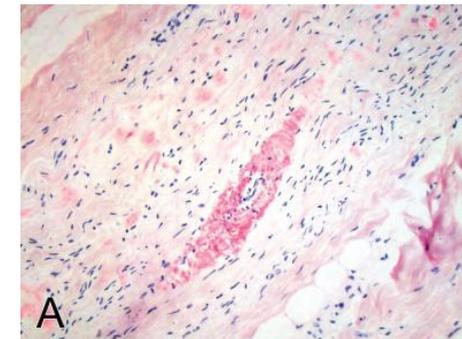
➤ hATTR Amyloidosis manifests with a variety of symptoms



Data from Alnylam (<http://www.alnylam.com/patients/alnylam-act/>) Mora et al. *Neurology* 2018;90(15 Suppl.):P1.448

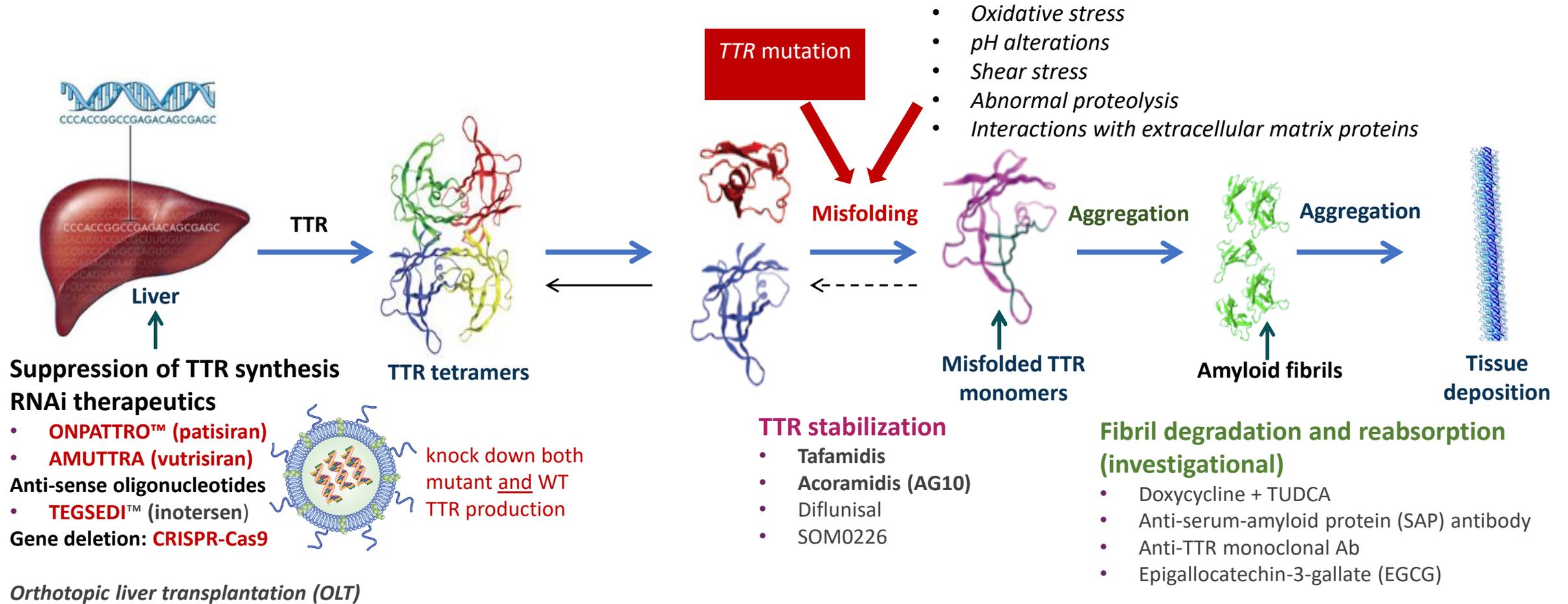


Rowcenzio DM, et al. *Hum Mutat* 2014;35:E2403-12



hATTR: molecular pathogenesis and therapeutic approaches

Misfolded mutant or wild-type TTR protein accumulates as amyloid deposits in nerves, heart, GI tract, and other tissues

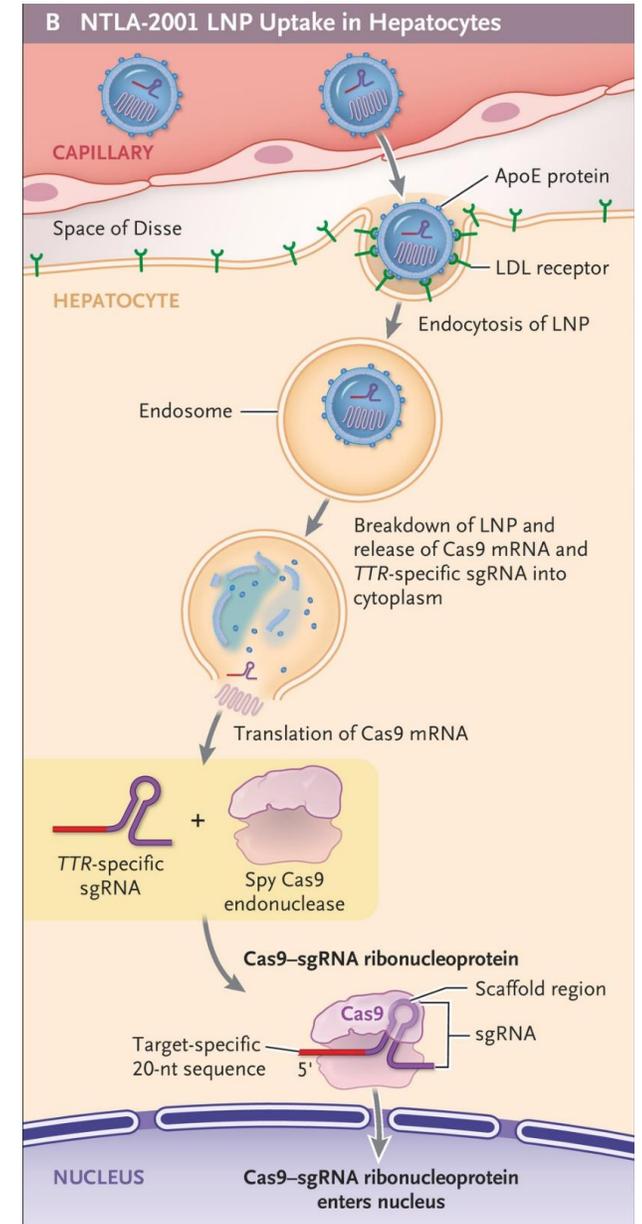
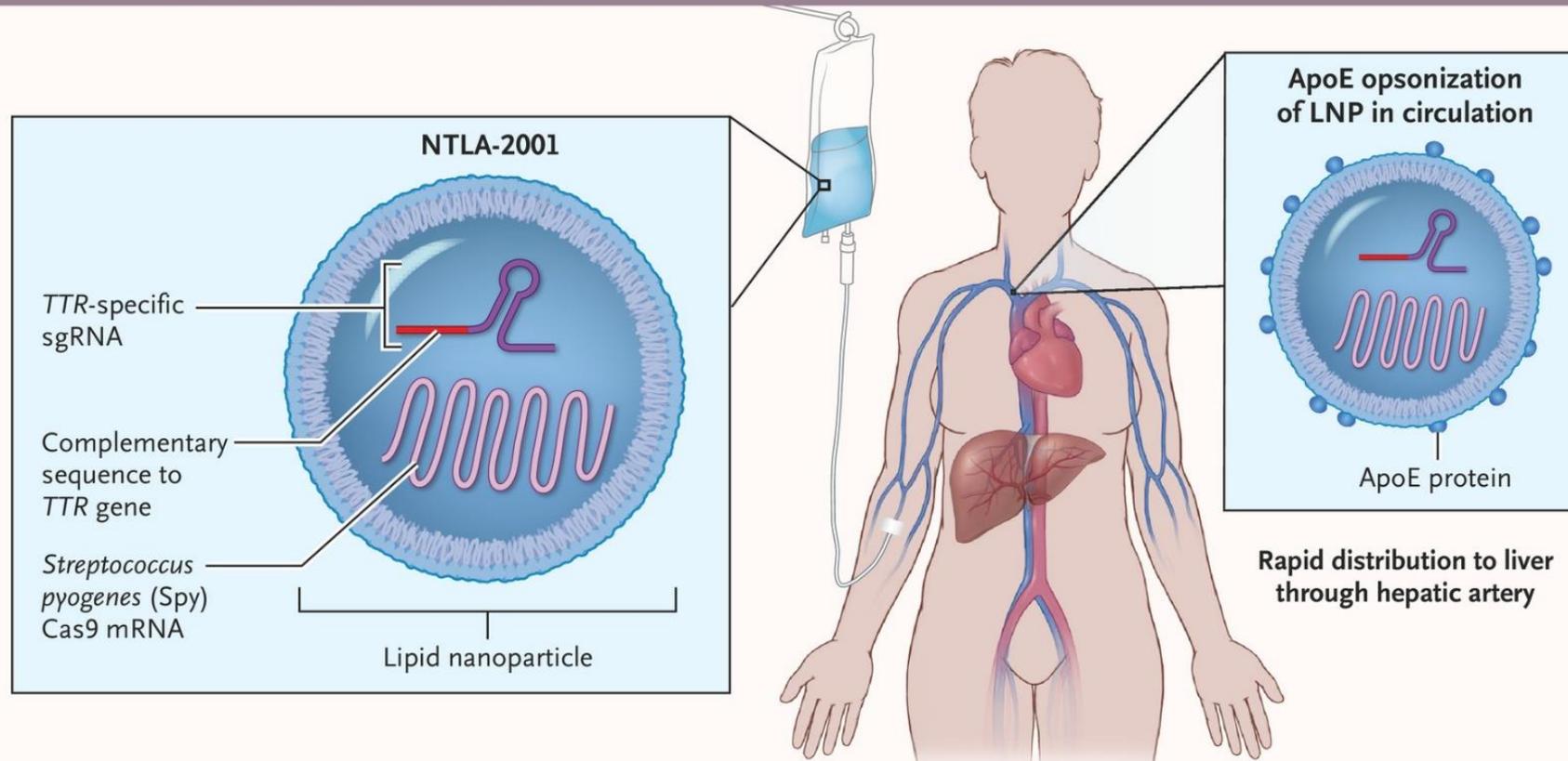


CRISPR-Cas9 In Vivo Gene Editing for Transthyretin Amyloidosis

Gillmore JD et al. DOI: 10.1056/NEJMoa2107454

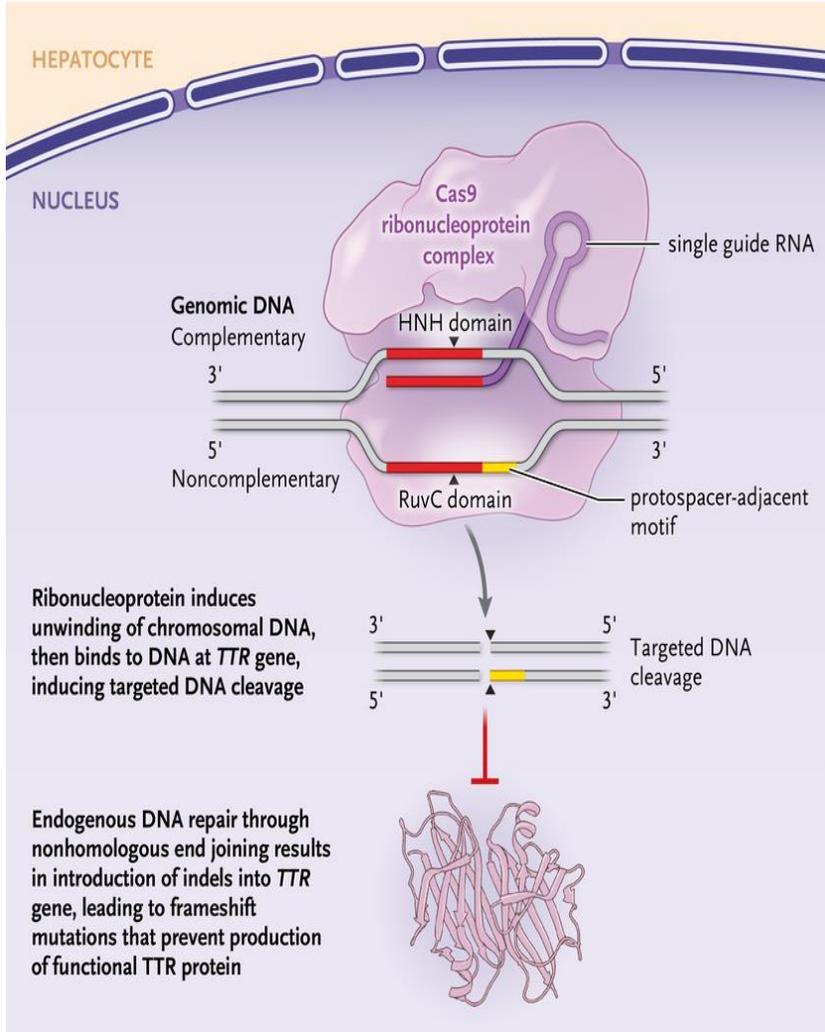
NTLA-2001: lipid nanoparticle (LNP) delivery system with liver tropism, carrying a single guide RNA (sgRNA) that targets human *TTR* and a human-codon-optimized mRNA sequence of *Streptococcus pyogenes* Cas9 protein

A Intravenous Infusion of NTLA-2001

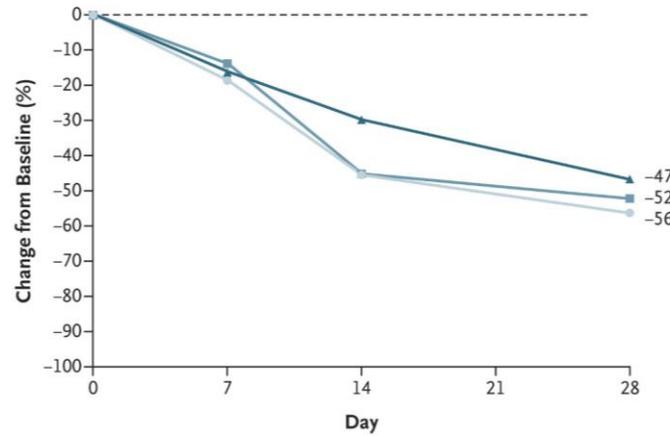


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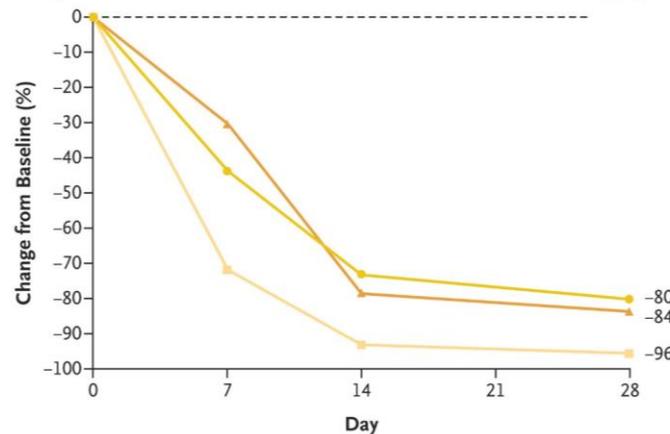
Gillmore JD et al. DOI: 10.1056/NEJMoa2107454



A Change in Serum TTR Concentration in Patients Who Received 0.1 mg/kg



B Change in Serum TTR Concentration in Patients Who Received 0.3 mg/kg



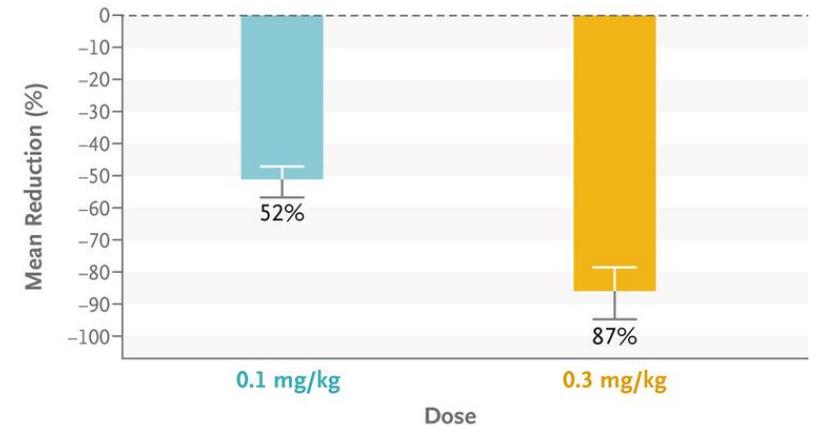
Phase I clinical study:

6 patients with hATTRv polyneuropathy

- 2 initial dose groups (n=3 of 0.1 mg/Kg and 0.3 mg/Kg)
- Ages: 53.5 years (46-64), 4 male, 2 female
- Polyneuropathy score 1 in all

Mean reduction from baseline in serum TTR protein concentration:

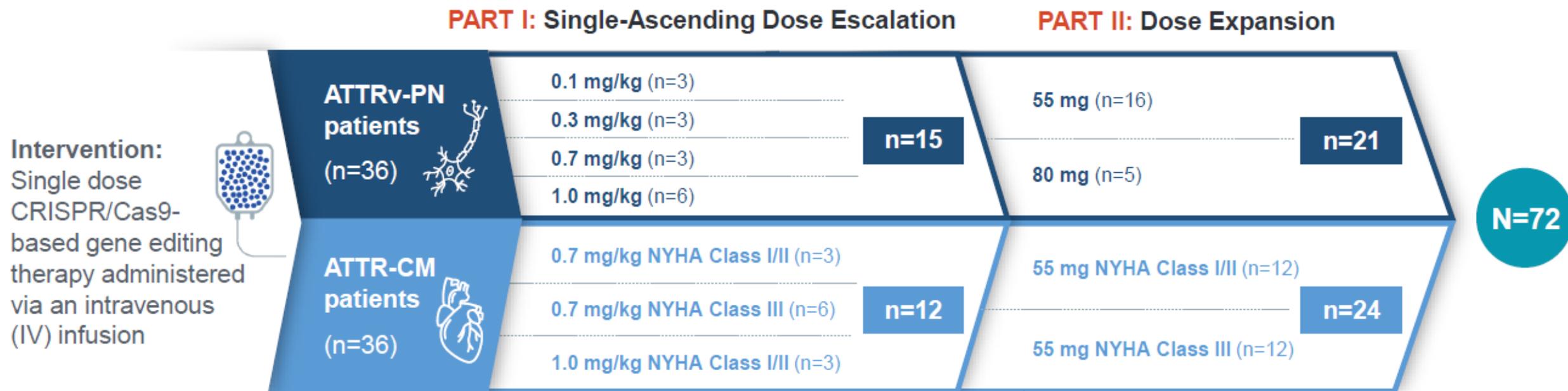
- 52% (47-56) in the 0.1 mg/Kg dose group
- 87% (80-96) in the 0.3 mg/Kg dose group



Two-part, open-label, multicenter study in adults with hereditary ATTR amyloidosis with polyneuropathy (ATTRv-PN) or ATTR amyloidosis with cardiomyopathy (ATTR-CM)

**Intellia
Therapeutics**

- **Ongoing Phase 1, two-Part: Open-label, Single Ascending Dose (Part 1) and Open-label, Single Dose Expansion (Part 2) Study to Evaluate Safety, Tolerability, Pharmacokinetics, and Pharmacodynamics of NTLA-2001 in Patients with ATTRv-PN and Patients with Cardiomyopathy (ATTR-CM)**
- **72 patients already treated with 4 dose-escalation cohorts**
- **expected completion in 2025**



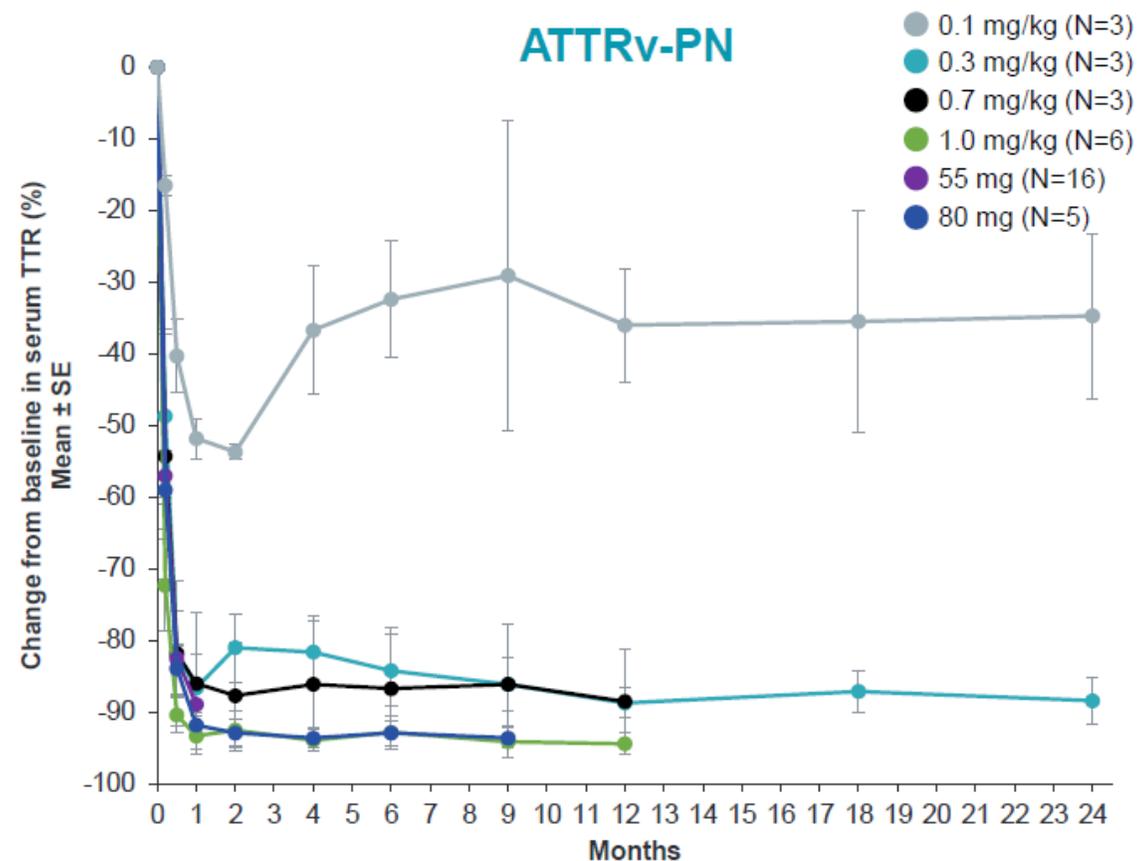
Presented by *Gillmore et al.* Nov 2023

CRISPR-Cas9 in vivo gene editing for ATTRv

Patient demographics and characteristics (PN arm)

Characteristic		PN Patients (N=36)
Age, years	Median (min, max)	61 (19, 75)
Sex, n (%)	Male	26 (72)
Weight, kg	Median (min, max)	77 (55, 117)
TTR genotype, n (%)	p.V50M	11 (31)
	p.V142I	1 (3)
	p.T80A	7 (19)
	p.S97Y	7 (19)
	p.E62D	4 (11)
	Other	6 (17)
	WT	0
NYHA Class, n (%)	No diagnosis of heart failure	12 (33)
	I	19 (53)
	II	5 (14)
	III	0
NT-proBNP, ng/L	Median (min, max)	127 (<50, 1878)

- NTLA-2001 led to **sustained reduction of TTR levels** in all patients and dose groups – but weaker reduction in the 0.1 mg/Kg group (data from 33 patients)



CRISPR-Cas9 in vivo gene editing for ATTRv: Safety and Side-Effect Profile

❑ From the published 2021 study

- Adverse events occurred during or after treatment in 3/ 6 patients, mild (grade 1) (Nausea, Headache, infusion related reaction)
- Increased D-Dimer levels observed 4-24 hours after infusion in 5 /6 patients, returned to normal by day 7
- Live function (ALS and AST levels) remained within normal limits

Gillmore et al. NEJM 2021

Potential for off-target gene editing with CRISPR-Cas systems?

- Therapeutic concentrations of NTLA-2001 showed no off-target mutagenesis mechanisms in primary human hepatocytes

➤ *Need to undergo long-term safety monitoring!*

❑ Preliminary safety data from the ongoing multicenter study

AE, Preferred Term, n (%)	Any Grade	Grade 1	Grade 2	Grade ≥3
Infusion-related reaction	25 (38)	10 (15)	14 (22)	1 (2)
Headache	12 (18)	12 (18)		
Diarrhea	11 (17)	10 (15)	1 (2)	
Back pain	7 (11)	7 (11)		
COVID-19 infection	6 (9)	5 (8)	1 (2)	
Cardiac failure	6 (9)	2 (3)	2 (3)	2 (3)
Upper respiratory tract infection	6 (9)	6 (9)		
AST increased	5 (8)	3 (5)	1 (2)	1 (2)
Dizziness	5 (8)	5 (8)		
Fatigue	5 (8)	5 (8)		
Muscle spasms	5 (8)	4 (6)	1 (2)	
Vision blurred	5 (8)	5 (8)		
Atrial flutter	4 (6)		1 (2)	3 (5)
Constipation	4 (6)	2 (3)	2 (3)	
Rash	4 (6)	4 (6)		

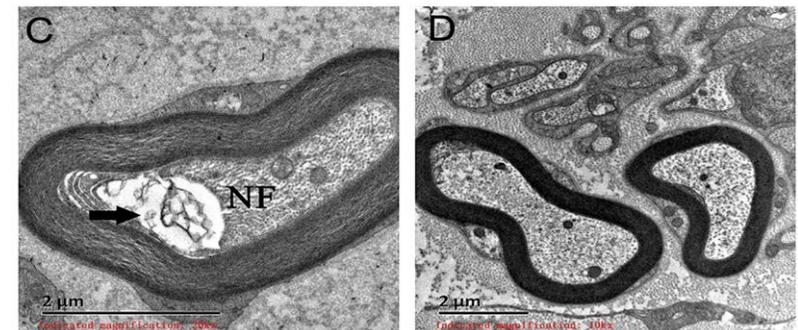
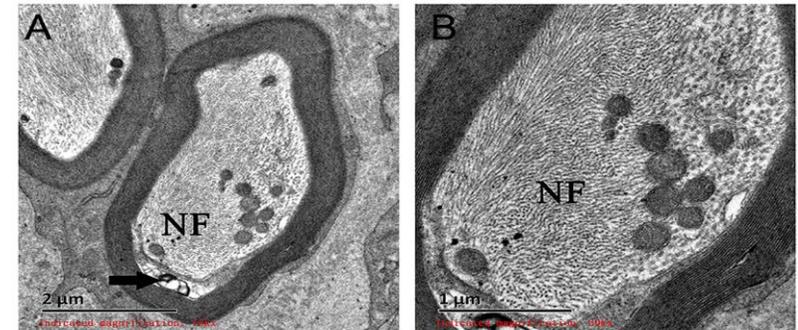
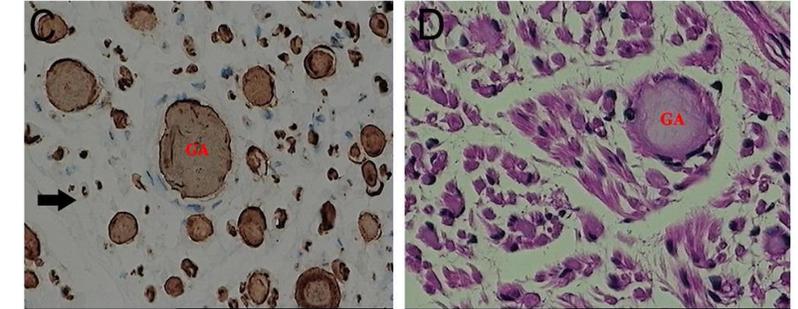
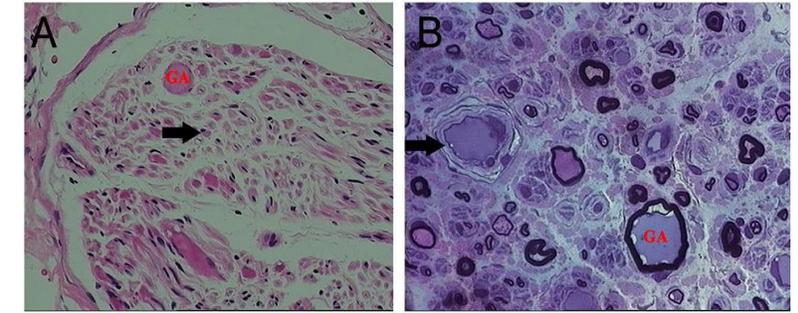
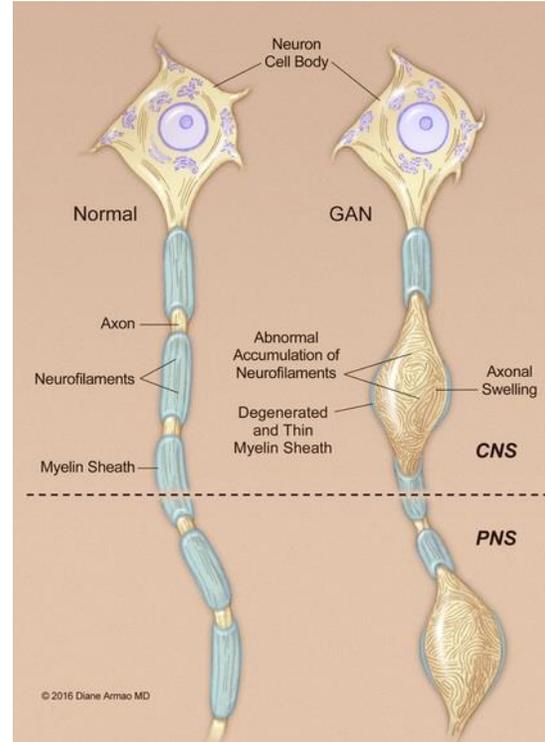
Presented by Gillmore et al. Nov 2023

Giant axonal neuropathy (GAN)

- Ultra-rare, autosomal recessive, progressive neurodegenerative disease with early childhood onset
- **Sensorimotor neuropathy** commonly progresses to affect **both the PNS and CNS**
- **Biallelic mutations in the GAN gene** located on 16q23.2, leading to loss of functional **gigaxonin**
- **Gigaxonin**: substrate-specific ubiquitin ligase adapter protein → regulates intermediate filament turnover

Loss of gigaxonin function

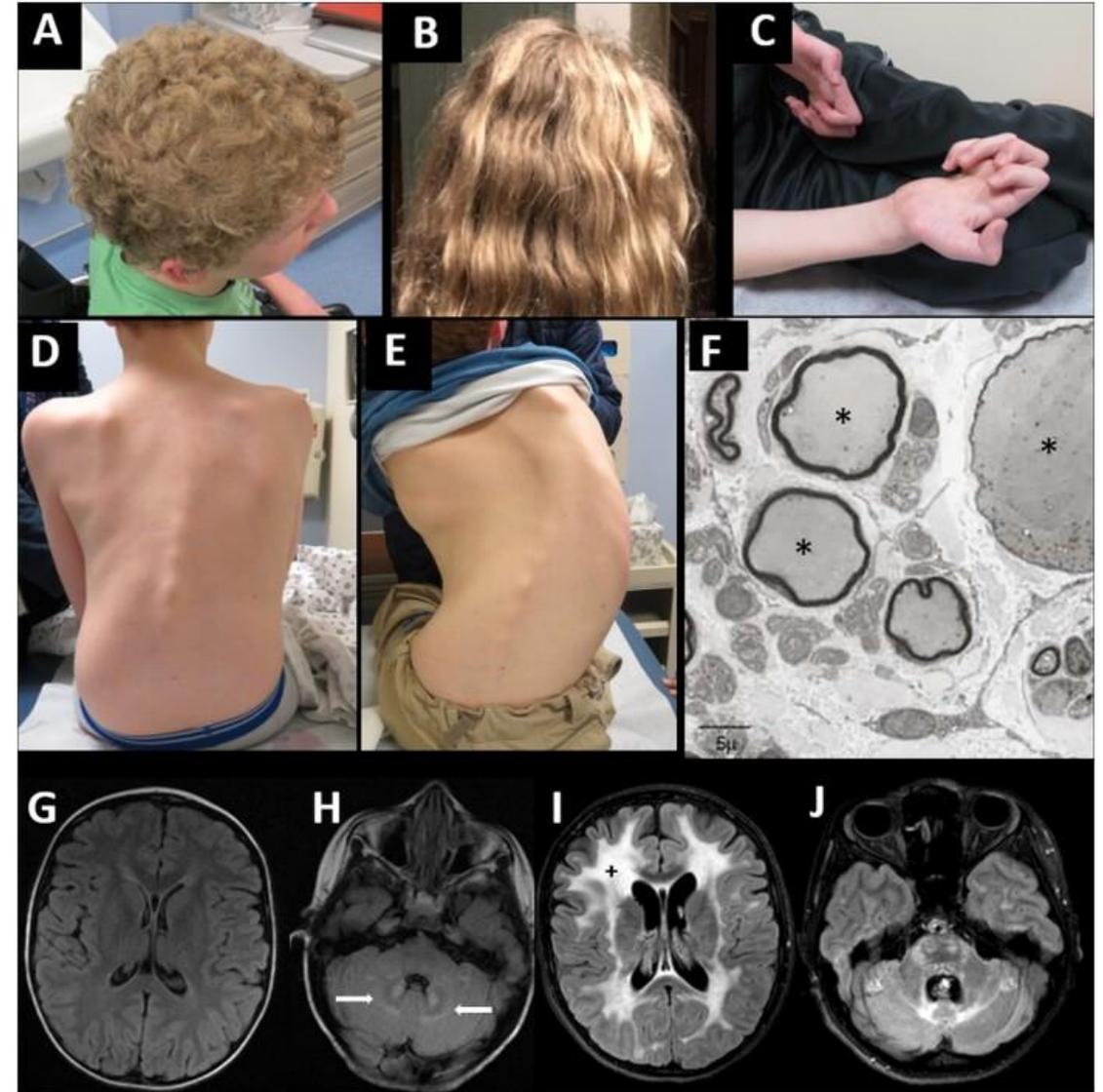
- **progressive accumulation of native intermediate filaments** in endothelial cells, skin fibroblasts, muscle fibres, Schwann cells, astrocytes and neurons
- **'Giant' axons or axonal swellings, within peripheral nerves, and in the CNS:** cerebral and cerebellar white matter, middle cerebellar peduncles, brainstem tegmentum, corticospinal tracts and posterior column (Asbury AK et al., 1972)



GAN clinical phenotype

Bharucha-Goebel et al Brain 2021: largest GAN cohort of 45 GAN patients (ages 3-21)

- **Mean age at symptom onset: 2.9 years**
- **PNS: Distal muscle weakness and sensory loss**
- Progressive **gait ataxia** in the ambulant individuals
- Age at **loss of unassisted ambulation: 8.3 years**
- **CNS:** Visual loss (22/45; 49%), dysarthria (19/45; 42%), urinary hesitancy or incontinence 13/45; 28%), precocious puberty (11/45; 24%), illness associated hypothermia (5/45; 11%) and sleep apnoea (14/45; 31%)
- **Seizures, vertigo, nystagmus, impaired cognitive development**
- Characteristic **dull, curly or tightly curled (frizzy) hair**
- **Gastrointestinal:** dysphagia (14/45; 31%), constipation (18/45; 45%); lactose intolerance (18/45; 40%)



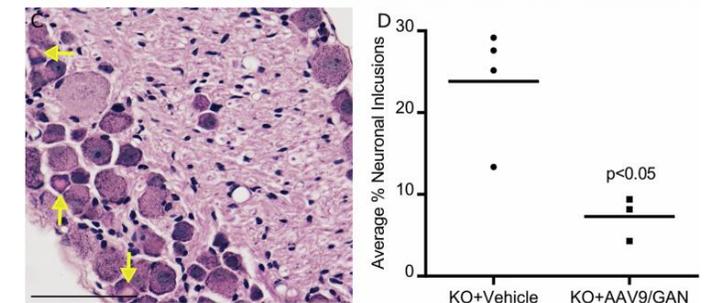
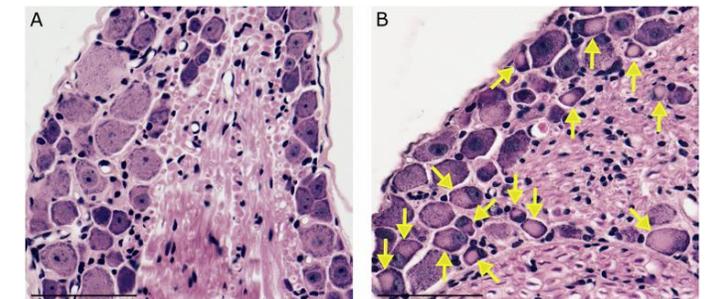
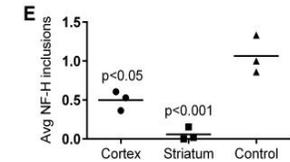
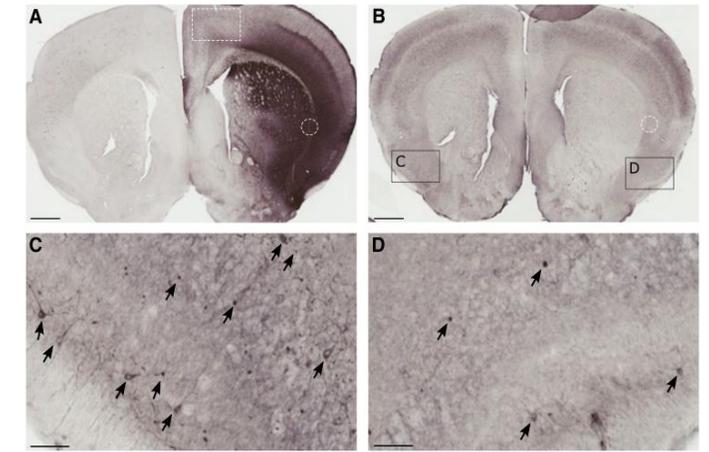
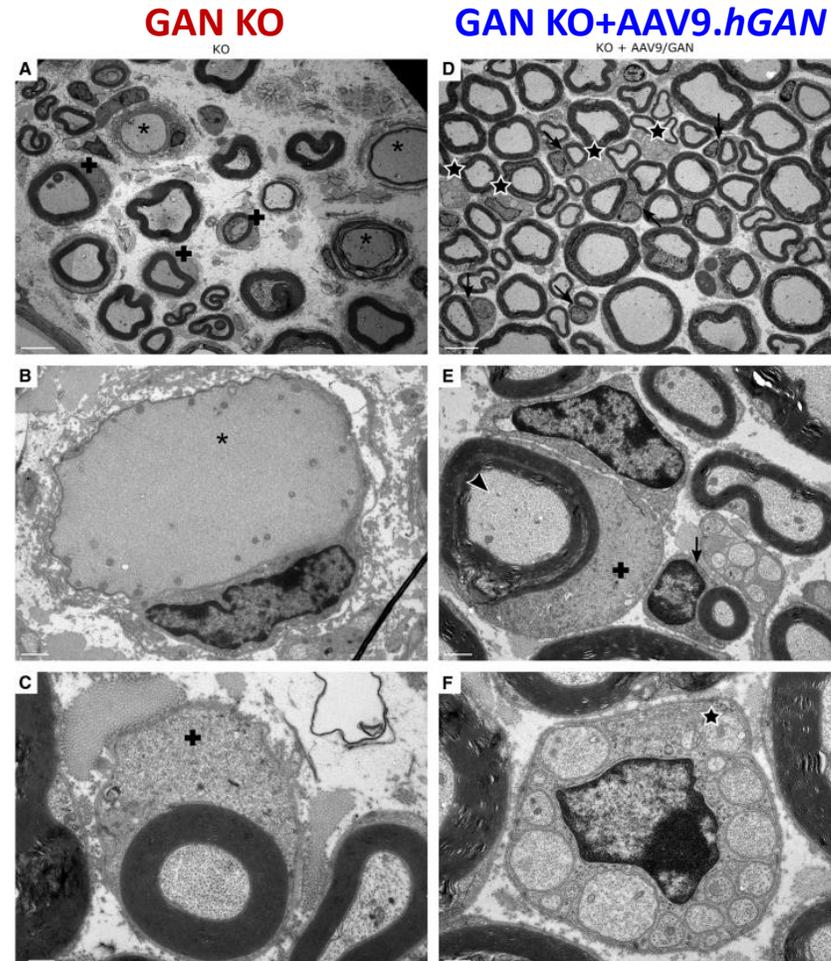
- Increased T2 signal abnormalities within cerebellar white matter surrounding the dentate nucleus of the cerebellum
- Cortical and spinal cord atrophy in more advanced disease severity

Development of GAN gene therapy



- AAV/JeT-GAN restored normal configuration of IFs in patient fibroblasts and in GAN KO mice
- IT delivery in aged GAN KO mice → preserved sciatic nerve ultrastructure
- → reduced neuronal IF accumulations
- → improved motor function
- Sustained wild-type gigaxonin expression in PNS and CNS for at least 1 year

Bailey et al., 2018



Clinical translation of GAN gene therapy

➤ **scAAV9-JeT-GAN** intrathecal gene therapy was well-tolerated and seemed to slow disease progression in GAN patients: [Data from Phase 1/2, first-in-human, single-site trial \(NCT02362438\)](#)

- **14 participants** with GAN dosed IT with either 3.5×10^{13} , 1.2×10^{14} , 1.8×10^{14} , or 3.5×10^{14} vg
- Corticosteroid **immunomodulation**, plus T-cell targeted immune modulation in biallelic null variants patients (n = 4)
- **Primary endpoints: safety and change in motor function measure (MFM-32) score** at 1 year compared to baseline

Safety:

- **TEAEs:** abnormal VC, URTI, UTI, acidosis, hyperglycemia, leukocytosis, thrombocytosis, CSF pleocytosis (clinically silent and self-limited), headache, intracranial hypertension
- **1 serious AE:** fever and vomiting requiring i.v. fluids likely related to treatment
- **2 deaths due to pulmonary complications: deemed related to GAN and not the gene therapy**
- **Anti-AAV9 antibodies:** peaked at 1:2,560 to 1:327,680 titers, 12 participants with increased T-cell interferon (IFN)- γ response (lower in participants receiving T-cell immune suppression)
- No dose-dependent side effects

Efficacy (1 year post treatment):

- **3 highest dose groups combined had a significantly slowed annual decline in MFM-32 score** ($P = .002$)
- **Nerve biopsies (n=11)** showed the presence of gene therapy in regenerative nerve fibers
- **Sensory nerve action potentials (SNAPs) persisted and/or re-emerged** compared to a decline in natural history
- **Increased regenerative clusters** in superficial sensory nerves

“Taysha Gene Therapies Halts Giant Axonal Neuropathy Drug Development on FDA Feedback”

! Lessons learned

- **Taysha Gene Therapies discontinued in Sep 2023 development of TSHA-120**, its gene therapy candidate for GAN, following feedback from the FDA on the company's intended registrational path.
- **“Due to challenges related to the feasibility of the study designs,”** (Taysha Chairman and CEO Sean Nolan)
- **Taysha submitted in 2022 available evidence from a Phase I/II trial of TSHA-120** and reviewed the data with the FDA
- **FDA provided feedback:** company needs to address **patient heterogeneity** in terms of disease progression in GAN and **concerns about using MFM32 as an endpoint**
- **Taysha subsequently submitted another analysis of data from a natural history and interventional trial comparing functional and biological measurements against a disease progression model**
- **FDA continued to recommend a randomized, double-blind, placebo-controlled trial to demonstrate efficacy** and suggested a **potential path for a single-arm trial with an external control group** and recommended **longer-term follow-up**
- **However, Taysha has determined that discontinuing development of the gene therapy will reduce operating expenses →** now focuses on its gene therapy trial for Rett syndrome.
- **Astellas Gene Therapies, which last year invested \$50 million into Taysha, has elected not to exercise its option to obtain an exclusive license to TSHA-120**

➤ **Careful clinical trial design!**

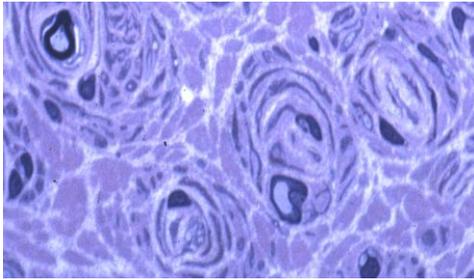
➤ **Well-designed Natural History Studies and carefully selected patients !**

➤ **Comprehensive, sensitive, regulatory acceptable outcome measures!**

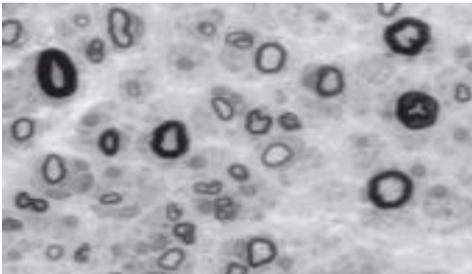
➤ **Challenging financial environment, prioritize resources!**

Genetic types and molecular mechanisms of CMT neuropathies

- Diverse cellular functions
- Many disease mechanisms
- Schwann cell expressed genes: → demyelinating CMT1, 4



- Neuronally expressed genes: → axonal CMT 2



Myelinating Schwann cell: CMT1, CMT4

Schwann cell cytoskeleton and linkage to extracellular matrix: *INF2, FGD4, PRX, FBLN5*

Transcription, mRNA processing: *EGR2, CTD1P1*

Mitochondria *GDAP1, HK1*

Endosomal sorting and cell signalling *LITAF/SIMPLE, SH3TC2, MTMR2, MTMR13, SBF1, FIG4, DNM2, NDRG1*

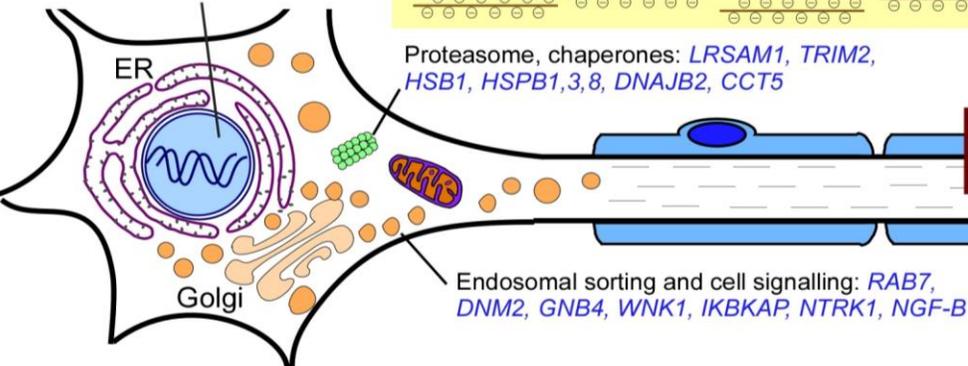
Compact myelin: *PMP22, MPZ*

Non-compact myelin: *GJB1*

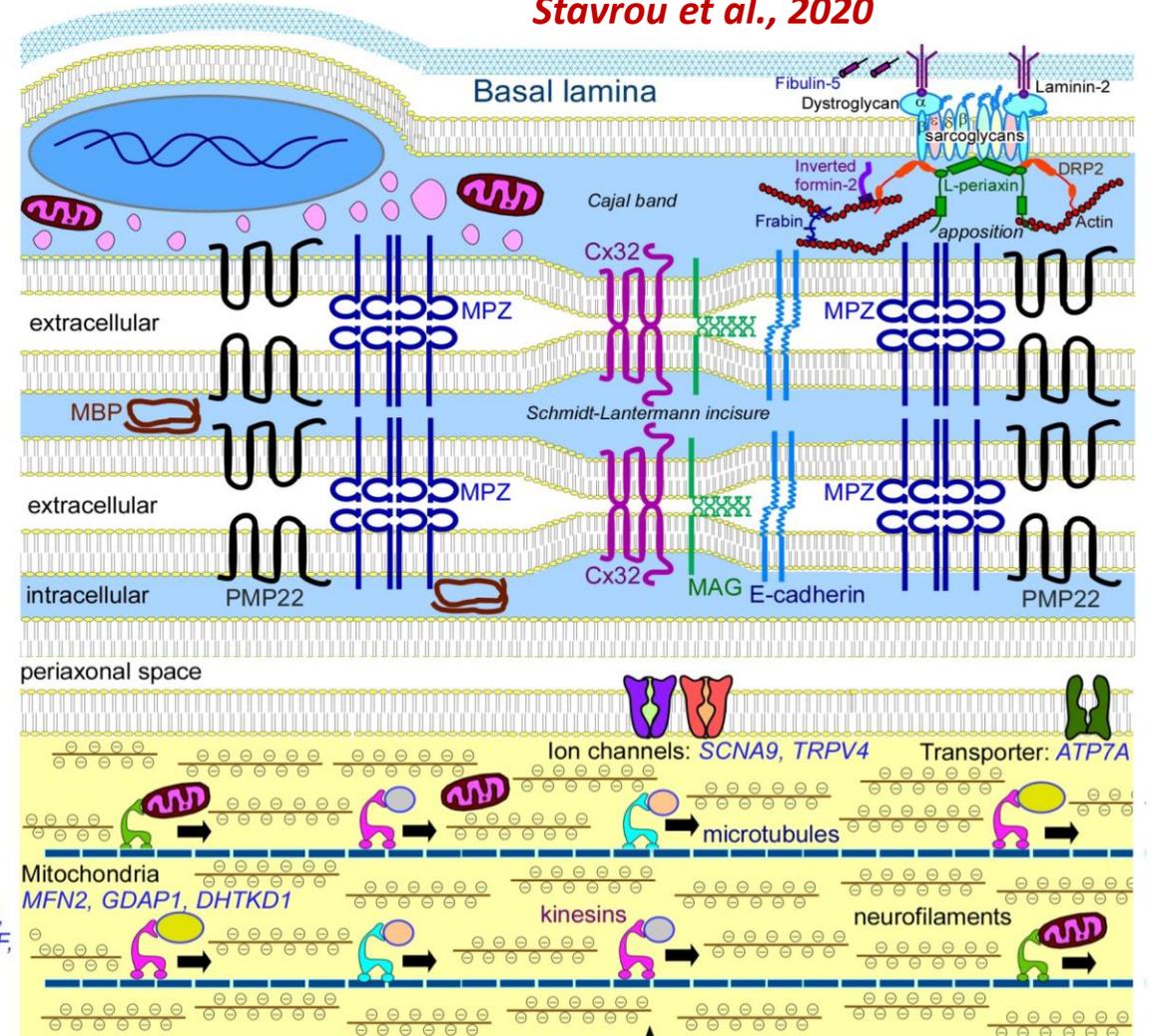
Neuron and axon: CMT2

Axonal transport: *NEFL, NEFH, KIF5A, KIF1A, DYNC1H1, HSPB1, BICD2, MYH14*

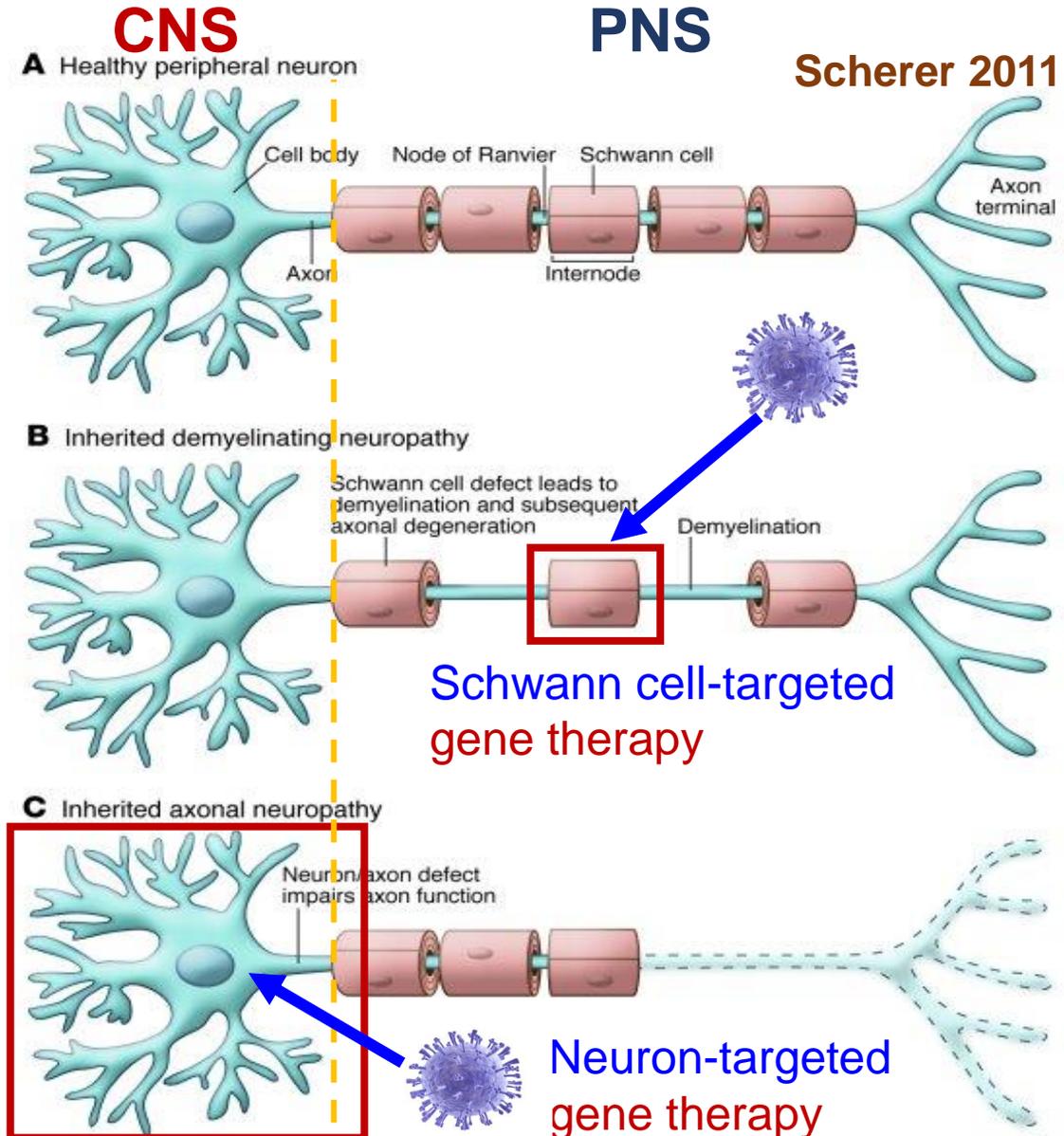
Nuclear envelope, mRNA processing: *LMNA, GARS, AARS, YARS, KARS, MARS, HARS, TGF, HINT1, PRPS1, IGHMBP2, DNMT1, MED25, PLEKHG5*



Stavrou et al., 2020



Planning gene therapies for CMT neuropathies



Gene therapies should:

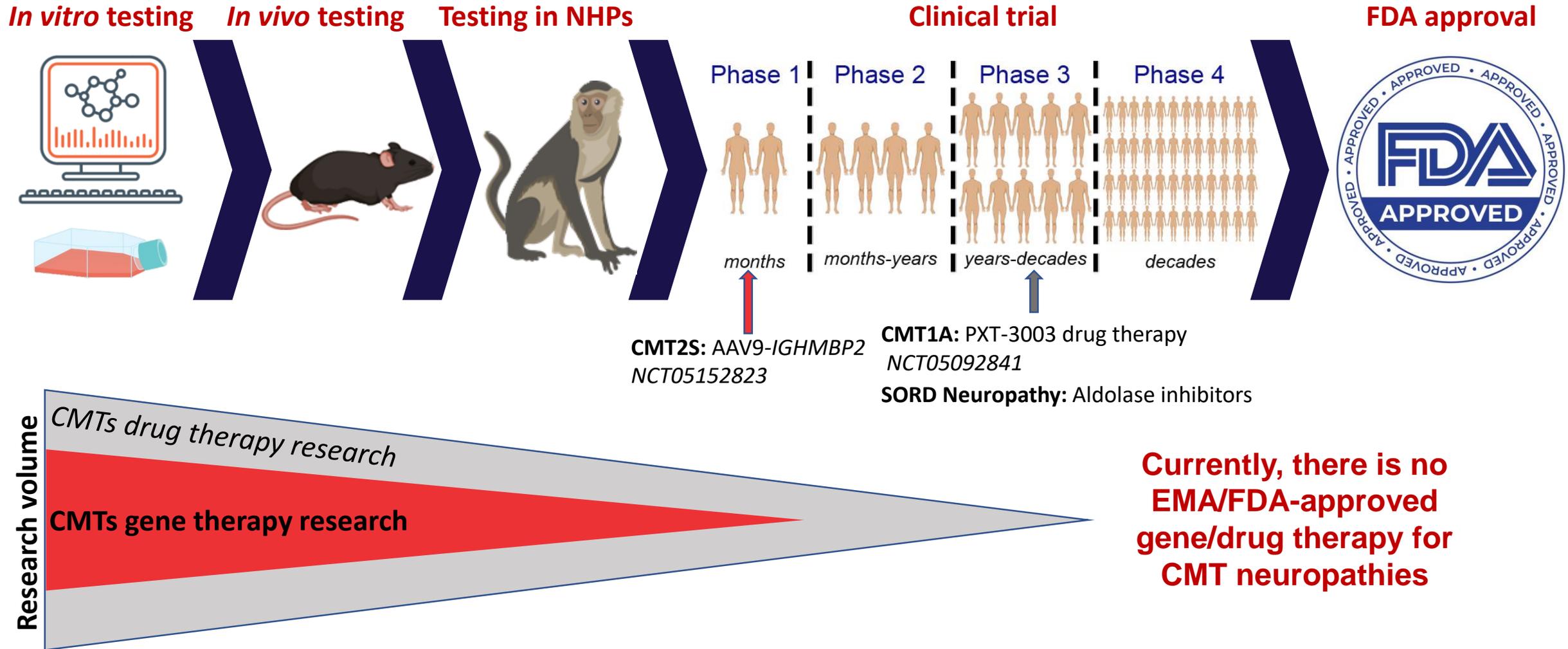
➤ address the **disease mechanism**:

- **Loss of function** → gene replacement
- **Gain of function** → gene silencing, allele-specific silencing, and/or editing

➤ be delivered to the **affected cell type**:

- Therapies for demyelinating CMT have to be **targeted to Schwann cells throughout the PNS**
- Therapies for axonal CMT neuropathies need to be **delivered to neurons** and their axons

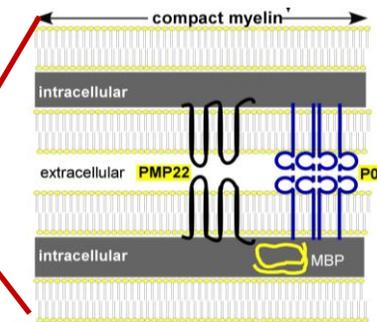
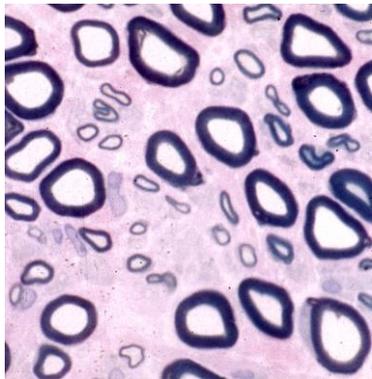
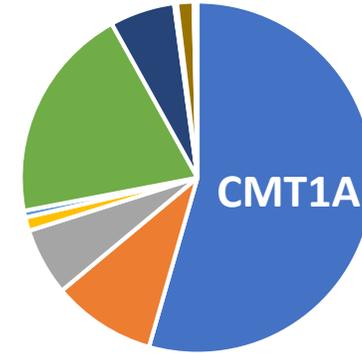
Long route toward translation of CMT gene therapies



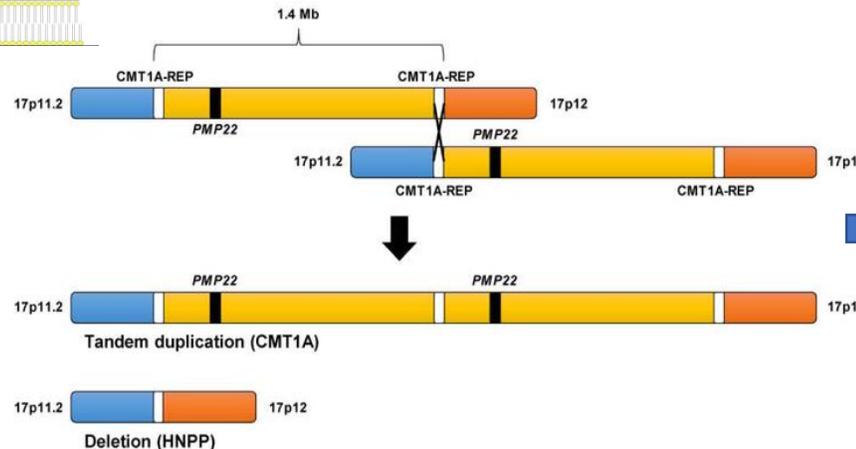
PMP22 duplication causes CMT1A

- **CMT1A** is the **most common** CMT type (~60 % of all CMT cases)
- Caused by peripheral myelin protein 22 (**PMP22**) **duplication**
- **Weakness and wasting** of distal muscles (feet > legs > hands), difficulty walking, sensory loss, impaired balance with **slow progression**, gradually increasing disability

CMT Subtypes by frequency



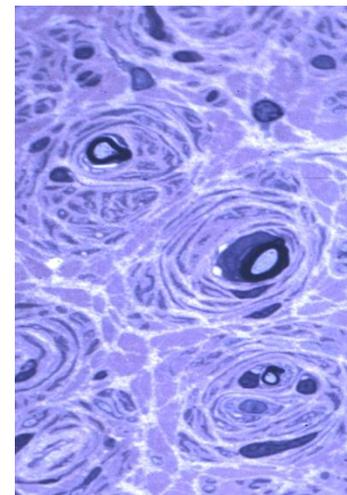
Unequal crossing over during meiosis → 1.4 Mb duplication on chrom. 17 p11.2



□ **PMP22** regulates Schwann cell growth and differentiation, myelin formation and maintenance

➤ **PMP22 overproduction** leads to accumulation, decreased proteasomal activity, ER stress, myelin dysregulation and demyelination

→ **Increased gene dosage effect!**

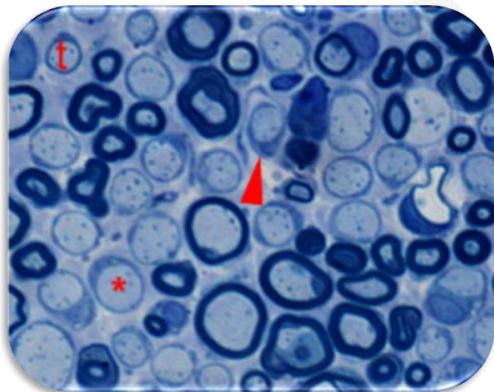


Gene therapies to reduce PMP22 levels

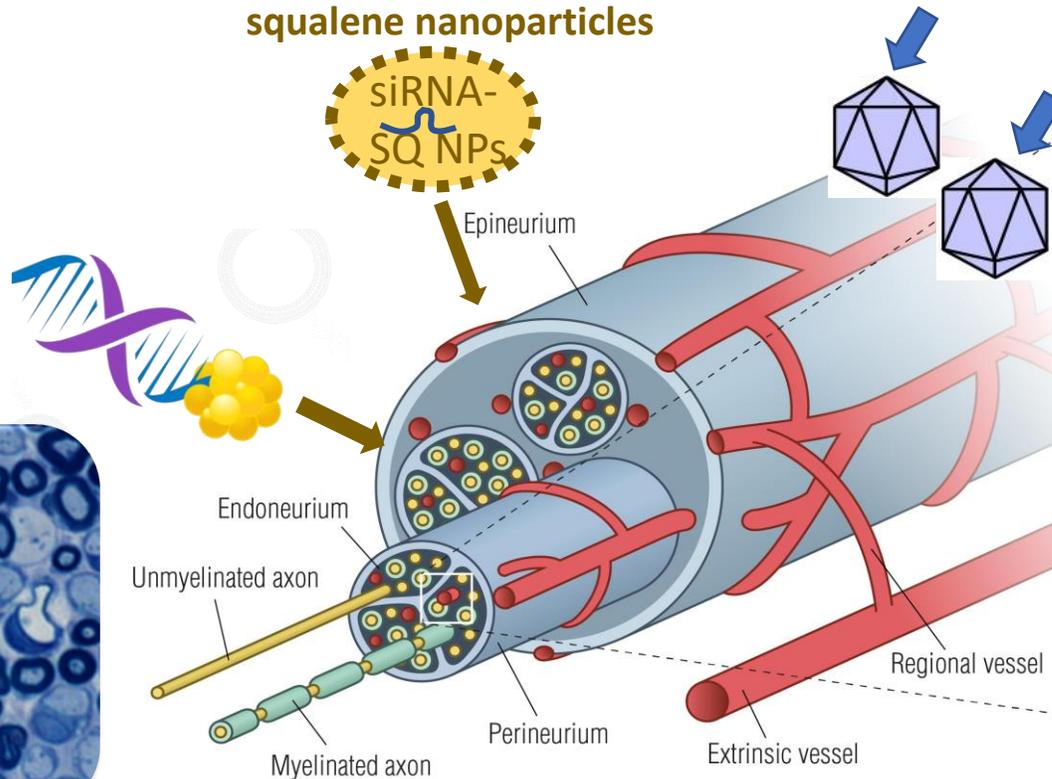
Non-viral methods

- Liposomes
- Nanoparticles
- ASOs

FALCON™ siRNA
(Fatty Acid Ligand
Conjugated
OligoNucleotide)-
DTx-Novartis



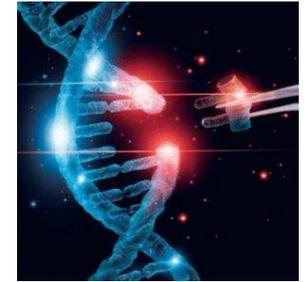
siRNA conjugated to
squalene nanoparticles



Viral vectors

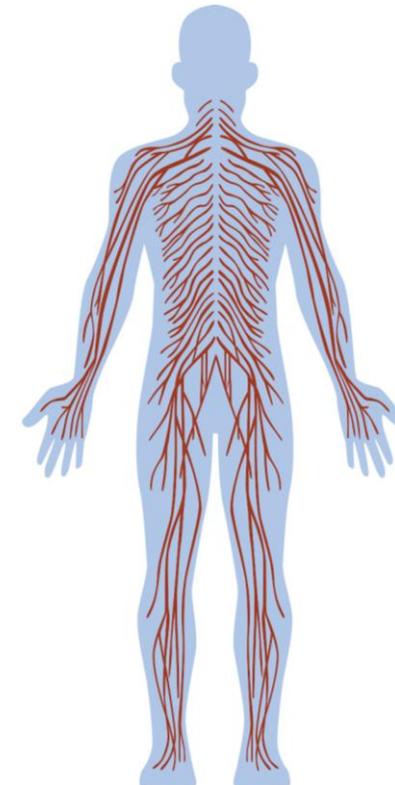
AAV vectors

- delivery of silencing molecules:
microRNAs, shRNAs, CRISPR-editing



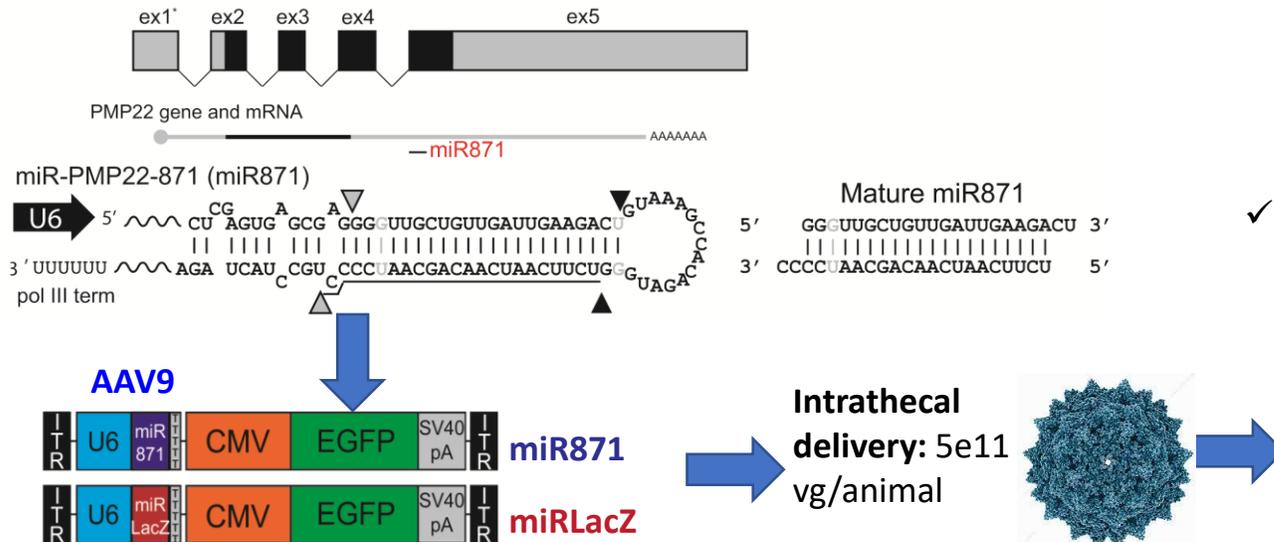
Delivery approaches for peripheral nerves

- **Subcutaneous** (ASOs, nanoparticles)
- **Intrathecal-preferable route for viral therapies**
- **Intravenous- high dose**
- **Intraneural** –restricted to injected nerve
- **Intramuscular-** for **trophic factors**

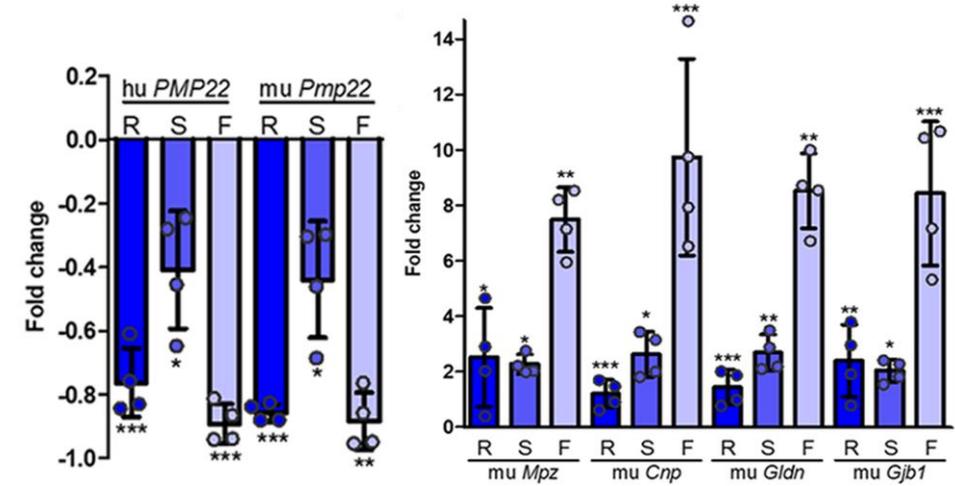


Development of gene silencing therapy for CMT1A

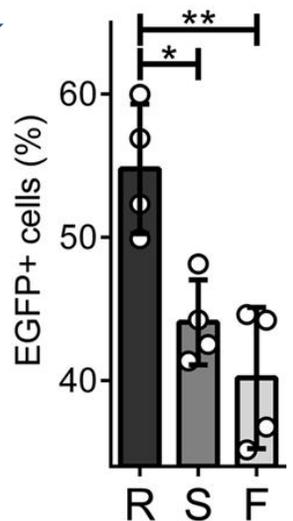
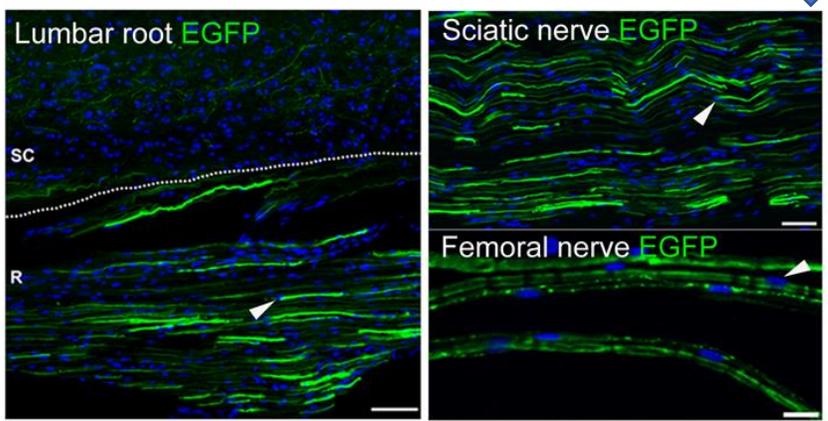
Stavrou et al., J Clin Inv 2022



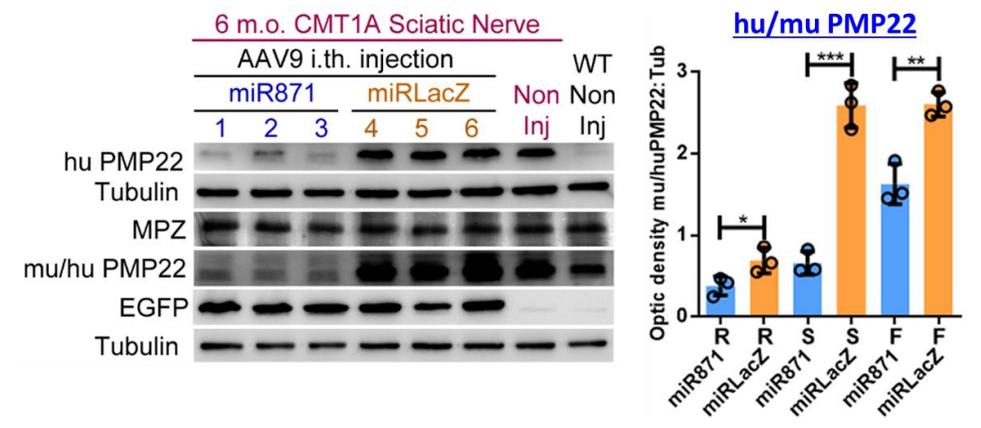
✓ Selective reduction of PMP22 mRNA in the CMT1A model



✓ Strong targeting of peripheral nerves and Schwann cell expression



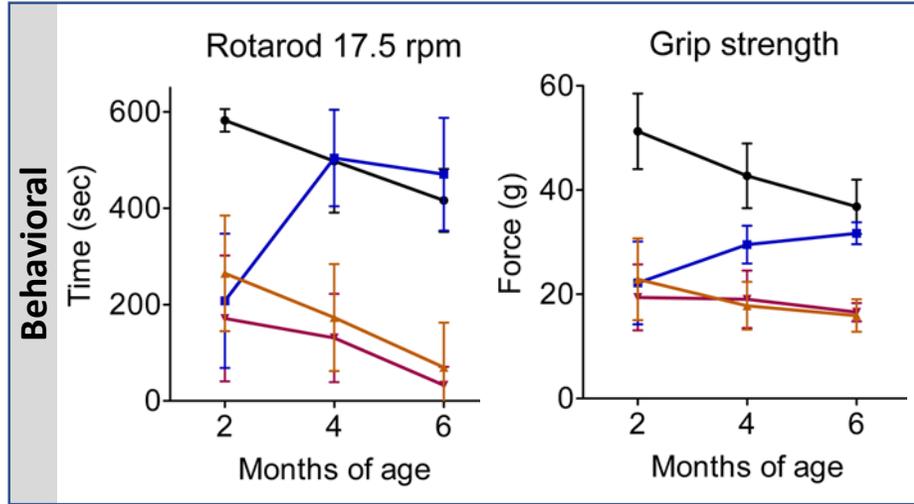
✓ Selective reduction of PMP22 protein levels



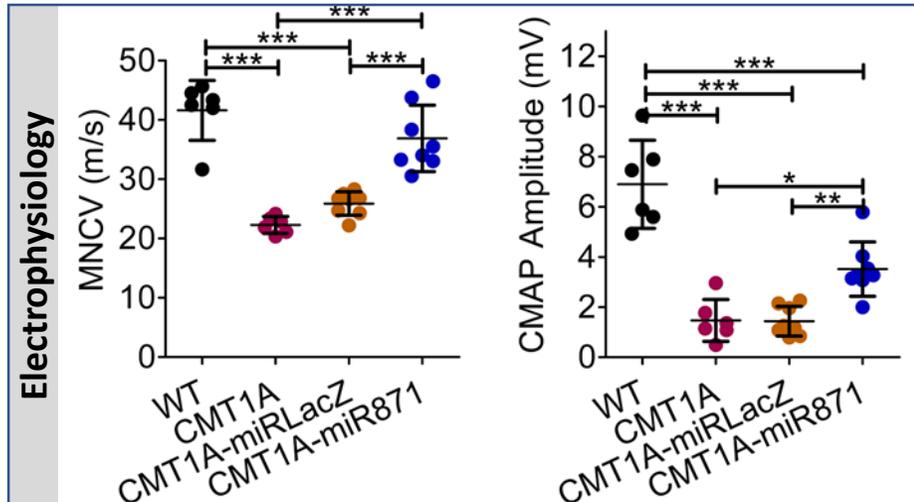
Development of gene silencing therapy for CMT1A

Stavrou et al., *J Clin Inv* 2022

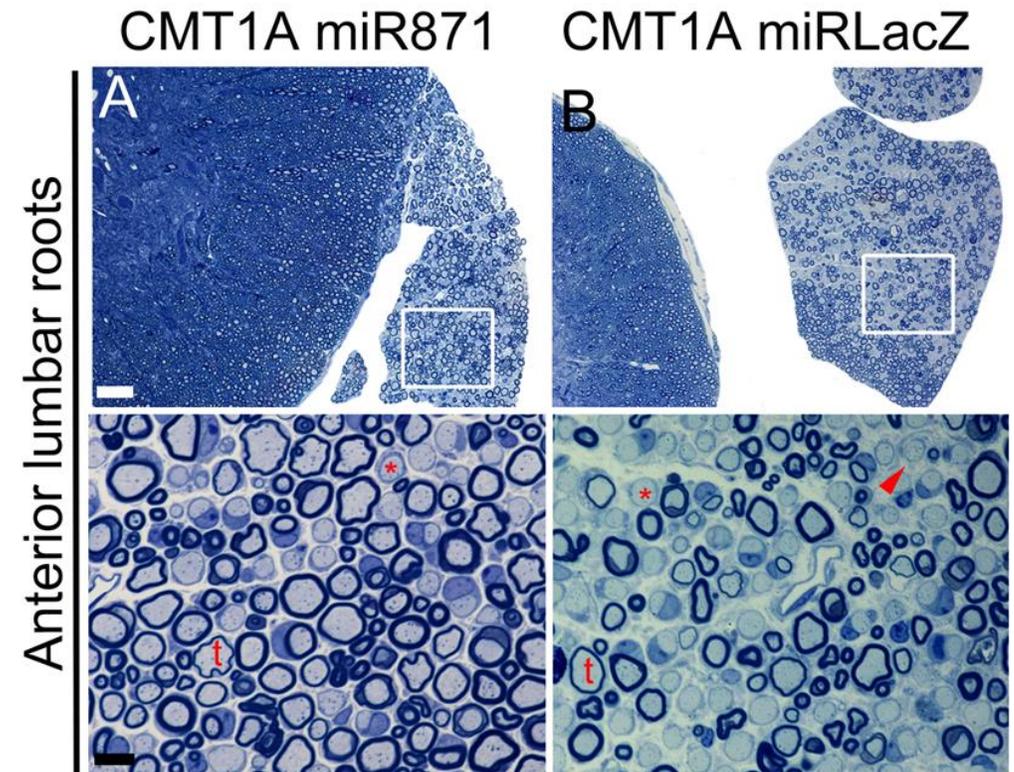
✓ Improved motor performance of the CMT1A model



✓ Improved nerve conduction velocities and CMAP



✓ Improved myelination, in lumbar roots and peripheral nerves of CMT1A mice after AAV9-miR871 treatment



Fatty Acid Ligand Conjugated Oligonucleotides (**siRNA**) targeting PMP22

DTx Pharma (acquired by Novartis in summer 2023)

FALCON technology overview:

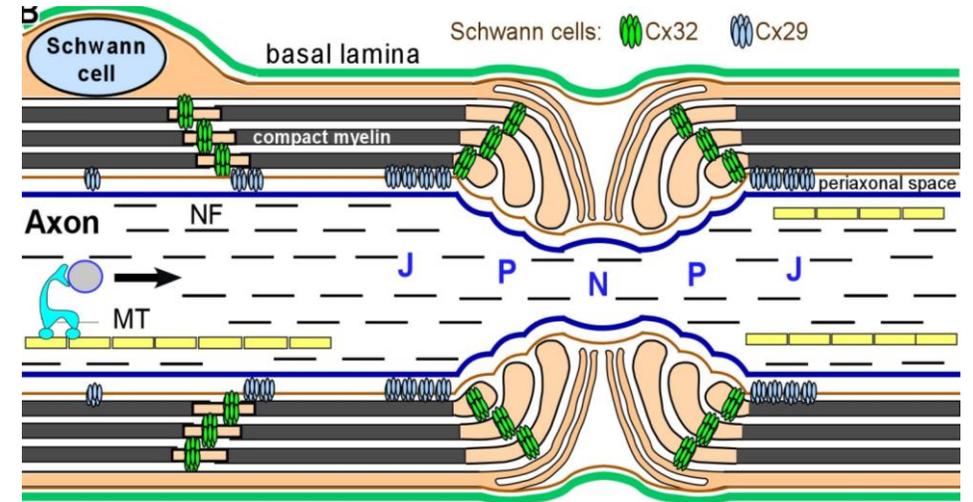
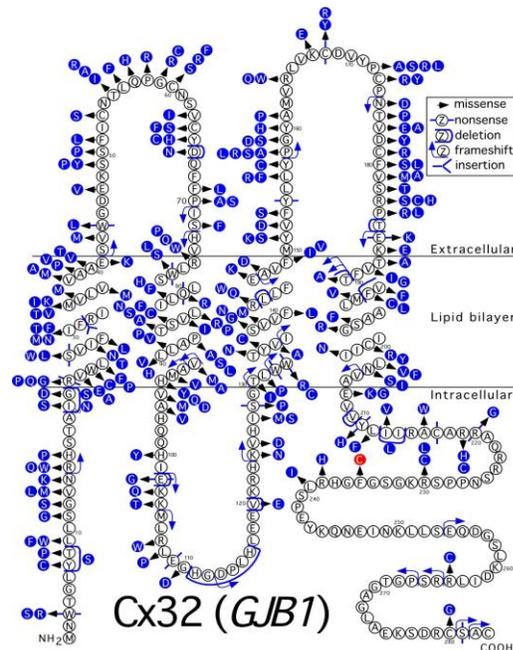
- Conjugates of naturally occurring fatty acids to siRNA's → **improve cellular uptake and biodistribution through binding to fatty acid receptors**
- **In vivo proof of concept** using the C3 mouse model (overexpressing human PMP22) of CMT1A
 - ✓ Remyelination of axons to normal levels
 - ✓ Improved muscle mass, grip strength, coordination and agility
 - ✓ “Reverse” multiple aspects of phenotype
- Completed **GLP Tox Studies and IND filing**
- Planning **first-in-human dosing in 2024**, with single and multi dose arms
- **Outcomes:** Safety, Biopsy, Electrophysiology (to evaluate remyelination)



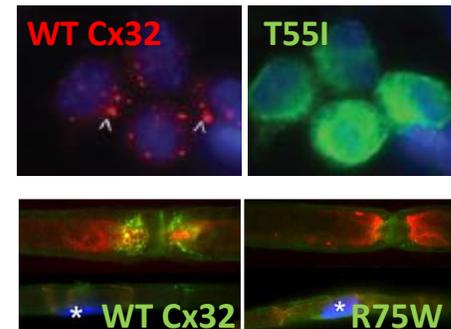
CMT1X phenotype and cause



- **CMT1X: 2nd most common** form of CMT (15-20%) – estimated prevalence of **1:25,000**
- Caused by mutations in the ***GJB1*** gene (on chromosome **Xq13.1**) encoding the **gap junction protein Connexin-32 (Cx32)**
- **Men affected earlier and more severely**
- **Symptoms usually start between 5 and 20 years in men:** difficulty running, ankle sprains, muscle atrophy
- Slowly progressive



- **Cx32 forms gap junction channels** through the myelin layers of myelinating Schwann cells in peripheral nerves
- **Communication pathway** transversing the myelin sheath **preserving homeostasis, axon support**



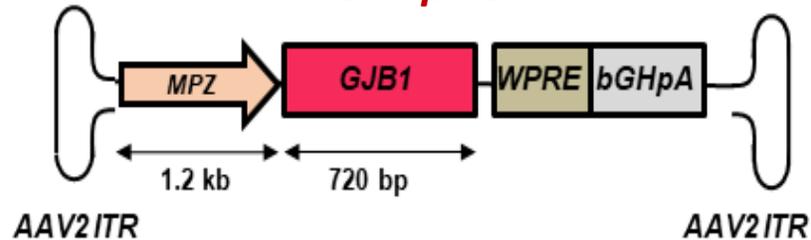
➤ **In CMT1X loss of Cx32 gap junction function leads to progressive degeneration of myelin and axon**

AAV-mediated *GJB1* gene replacement

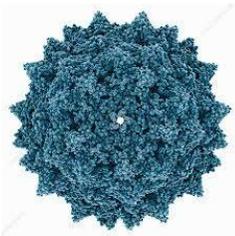
Kagiava et al., Sci Rep 2021;
Kagiava et al. Gene Therapy 2021;
Kagiava et al., Mol Ther Meth Clin Transl 2023

Therapeutic vector:

AAV9-Mpz.GJB1

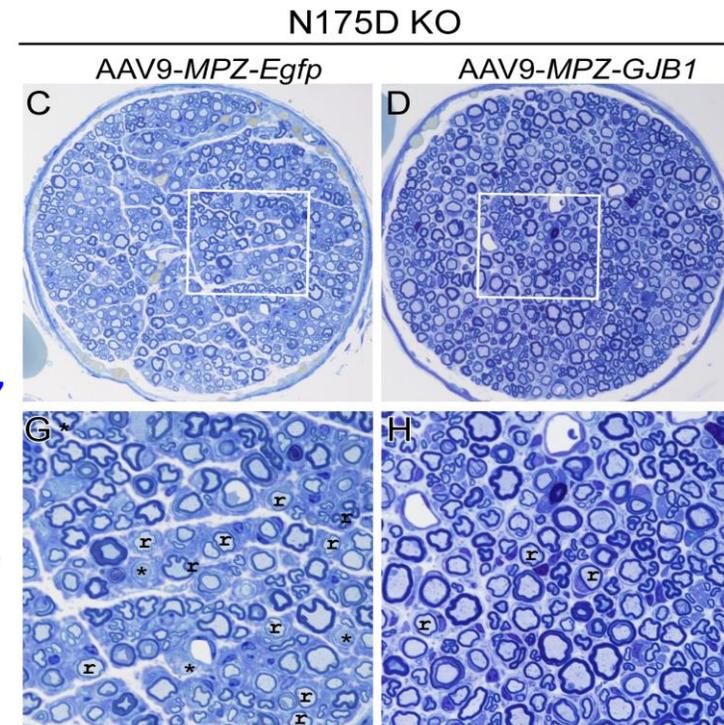
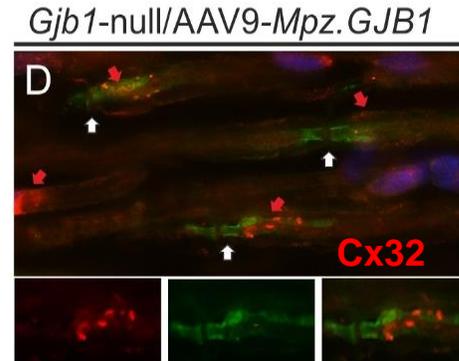


Myelin-specific myelin protein zero (Mpz) promoter



→ Lumbar intrathecal delivery in different models of CMT1X 2×10^{11} vg

- Adequate biodistribution to PNS tissues
- High percentage of Schwann cell-specific gene expression
- Pre-onset (2 month) and post-onset (6 month-old) treatment trial in CMT1X models provides therapeutic benefit
- Cx32 deficient (*Gjb1*-null)
- CMT1X transgenic (R75W and N175D):
 - ✓ **Functional improvement** (muscle strength, nerve conduction velocities)
 - ✓ **Improved myelination** (reduced demyelinated and thinly myelinated fibers)
 - ✓ **Reduced inflammation** (number of macrophages)



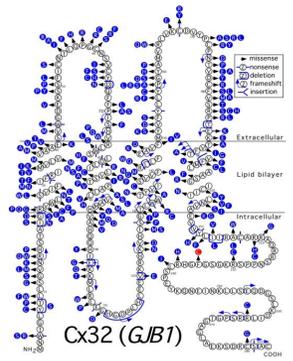
Route toward translation of CMT1X gene therapy

In vivo testing in disease models

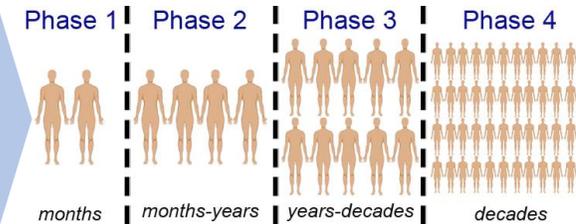
Testing in NHPs

Clinical trial

FDA approval



Disease cause and mechanisms



- POC studies completed → therapeutic benefit of IT AAV9 and AAVrh10 in different CMT1X models

- Dose-escalation and toxicity study completed successfully in the CMT1X model
- Further testing of leading capsids underway

- Non-human primate studies to assess **scale-up potential** and **safety** in **collaboration with industry**
- Completed 12/2023, final results Q2 2024

- Natural history studies
- Clinical trial readiness
- Biomarker validation (MRI, etc)

Gene therapy for SMA with respiratory distress type 1 (SMARD1)/ CMT2S

- **Recessive loss of function *IGHMBP2* mutations** → lower motor neuron loss → limb muscle atrophy and respiratory complications in infancy (SMARD1)
- **Partial loss of function** → milder phenotype with later onset: CMT2S
- **Dose-dependent gene replacement in the SMARD1 mouse model (*nmd*)**

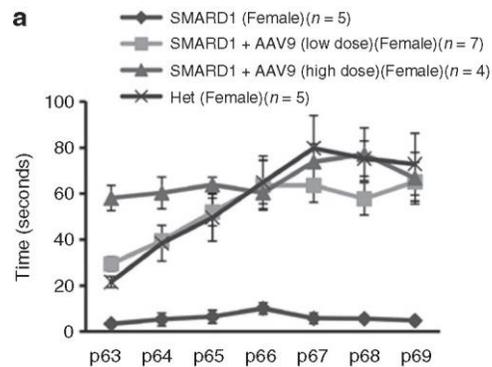
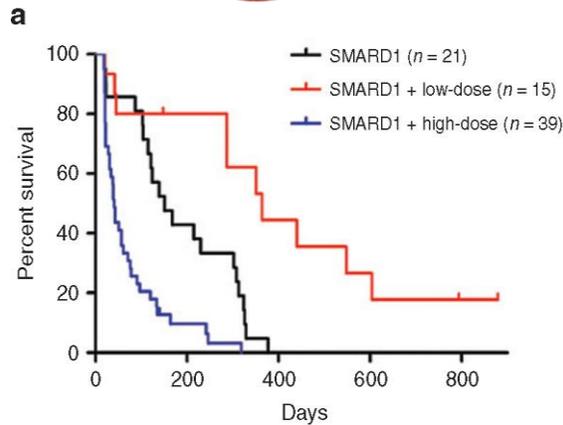
➤ **Clinical Trial** [NCT05152823](#) by **Alkyone Therapeutics** in children 2 mo-14 yr

Shababi et al., 2016, Mol Ther

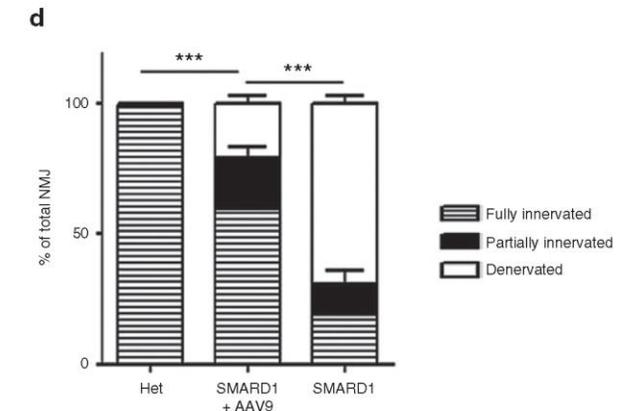
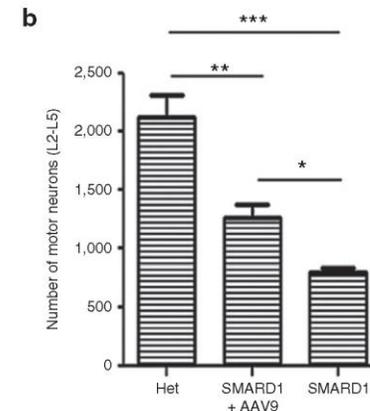
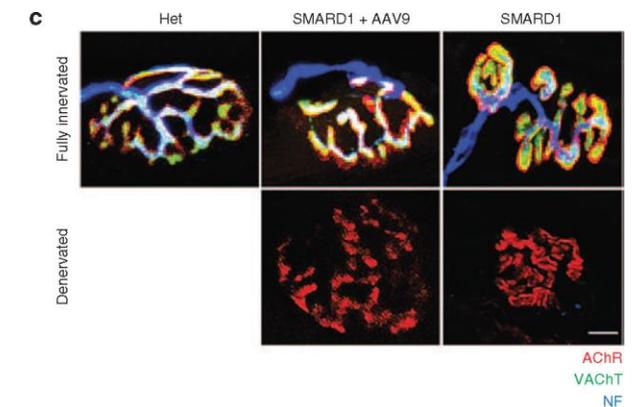
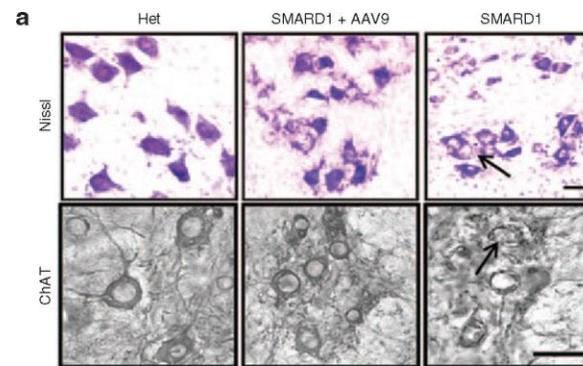
1.25-2.5e11 vg/animal



AAV9-CBA-IGHMBP2
post natal day 1-4



➤ **AAV9-IGHMBP2 rescued loss of motor neurons and NMJ innervation**

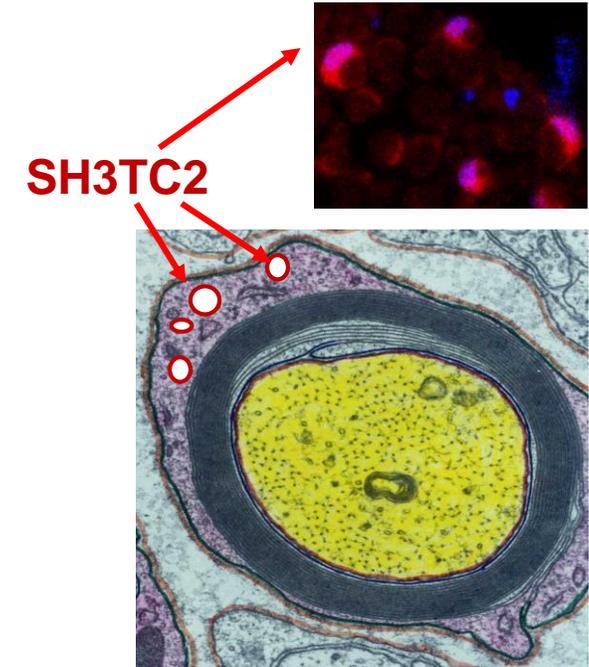


➤ **AAV9-IGHMBP2 *low but not high dose* increased survival and improved motor function**

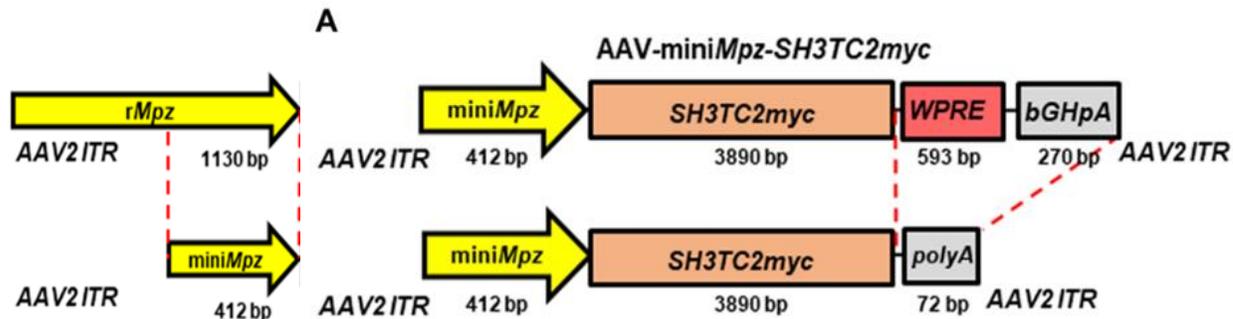


Gene therapy for CMT4C

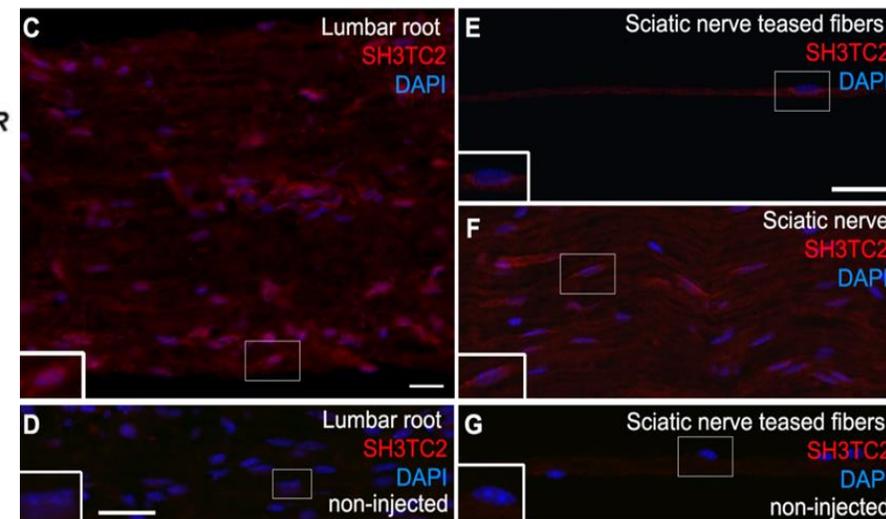
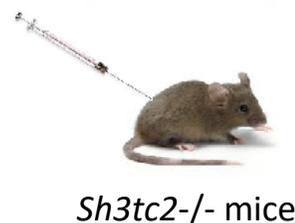
- **CMT4C: Severe, early onset demyelinating CMT neuropathy** with faster progression compared to CMT1 forms
- Caused by mutations in the **SH3TC2** gene -**most frequent CMT4 type: 3-4% of all non-CMT1A demyelinating CMT neuropathies**
- Autosomal recessive with **loss of function** mechanism
- **SH3TC2** protein is an **endocytic recycling compartment** protein specifically expressed by myelinating Schwann cells –interacts with Rab11 → **CMT4C mutations disrupt function**
- Early onset demyelination in *Sh3tc2*^{-/-} mice (*Arnaud et al., 2009*)



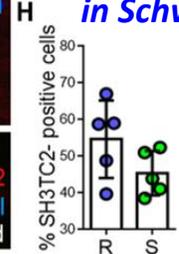
➤ Therapeutic vector generation for SH3TC2 gene replacement *Georgiou et al., Mol Ther 2023*



- **Lumbar intrathecal injection 20 μl (2E+11 vg) in *Sh3tc2*^{-/-} mice**



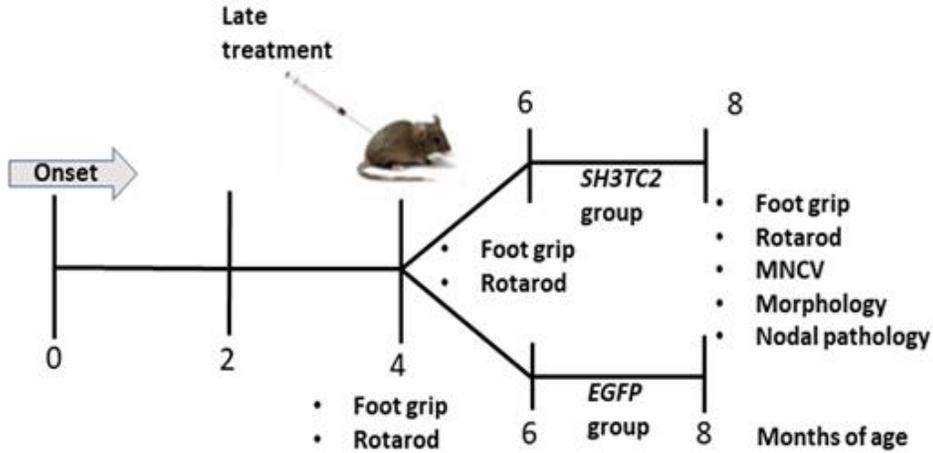
- **Sufficient vector biodistribution to the peripheral nerves**
- **High SH3TC2 expression rates in Schwann cells**



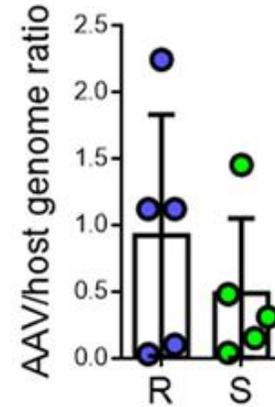
Functional improvement in treated *Sh3tc2*^{-/-} mice

Georgiou et al., Mol Ther 2023

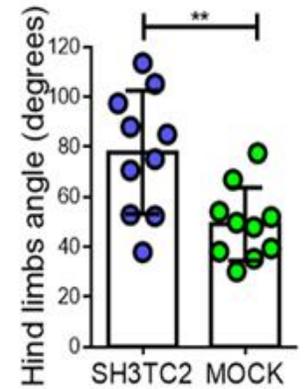
Study design



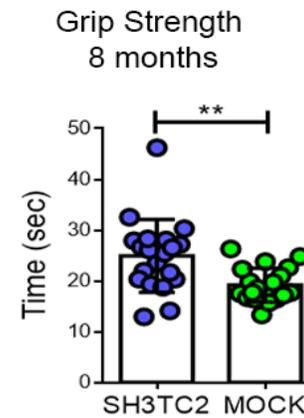
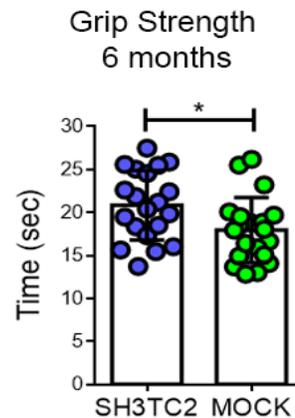
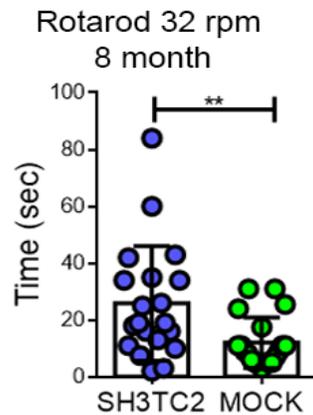
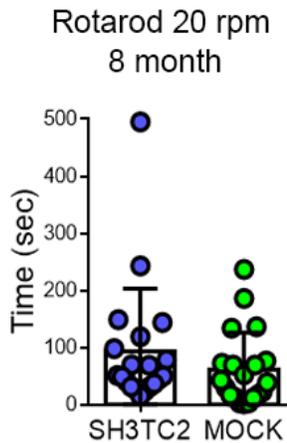
Biodistribution



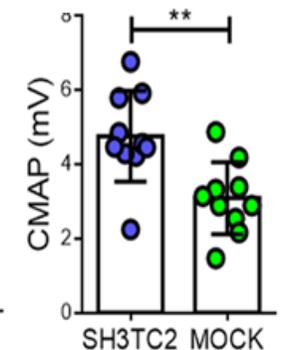
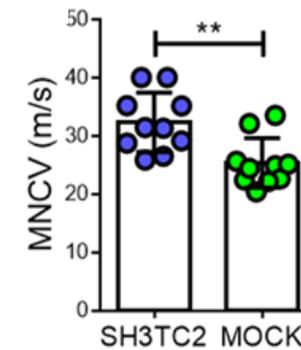
Behavioural analysis



Behavioural analysis

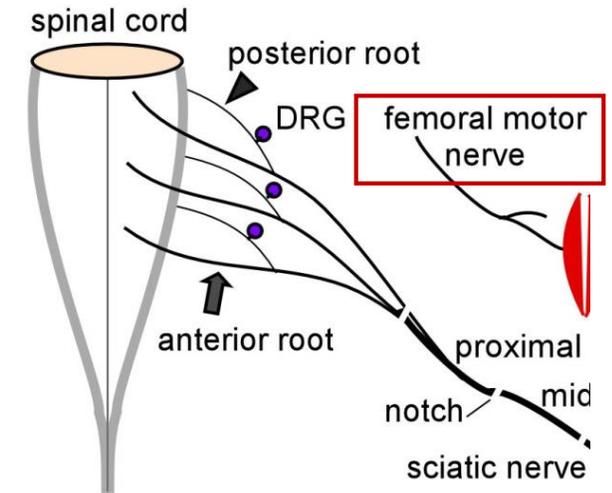
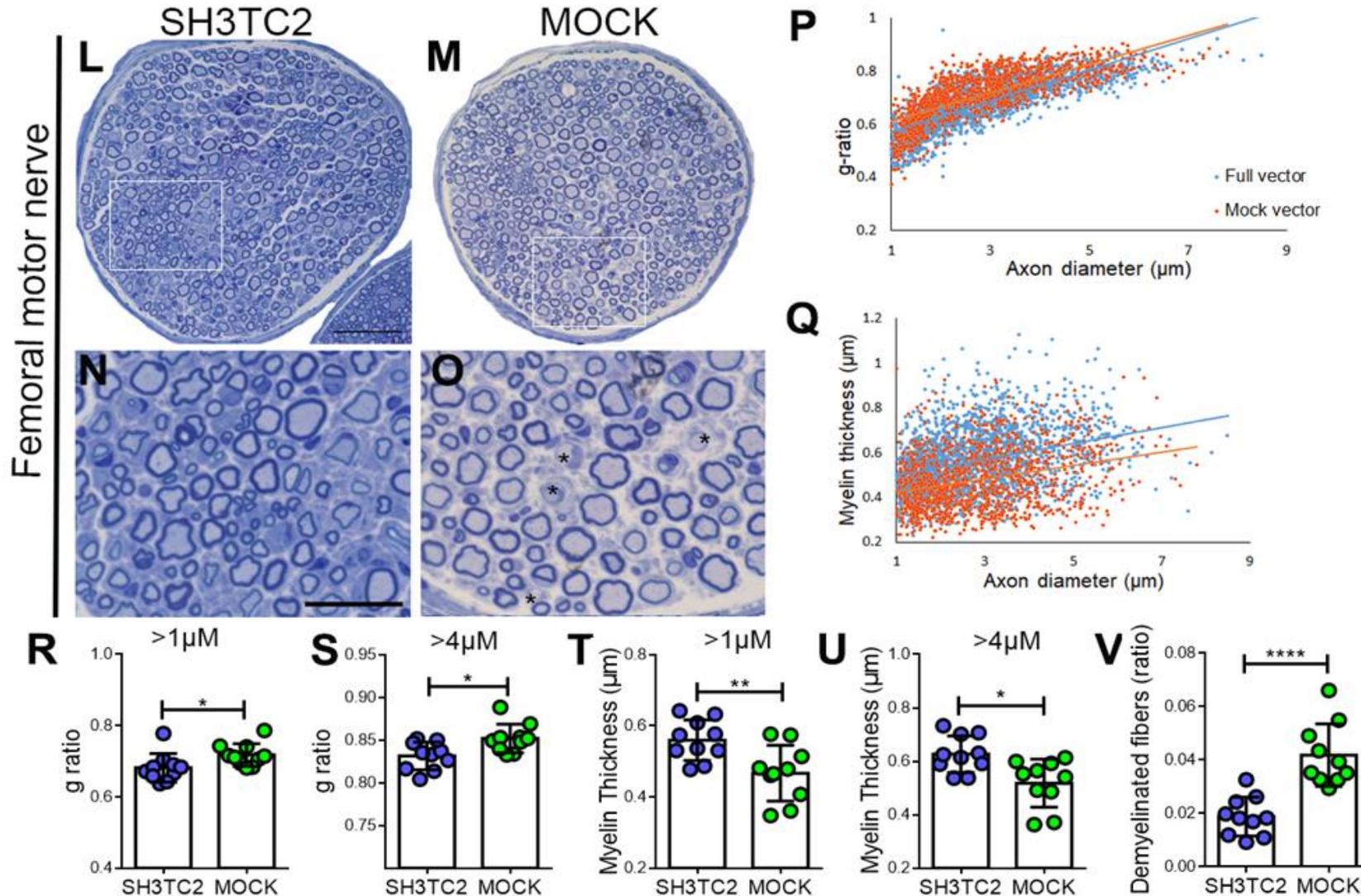


Electrophysiology



Improved myelination in treated *Sh3tc2*^{-/-} mice

Georgiou et al., *Mol Ther* 2023



➤ Similar results in lumbar motor roots and in sciatic nerves

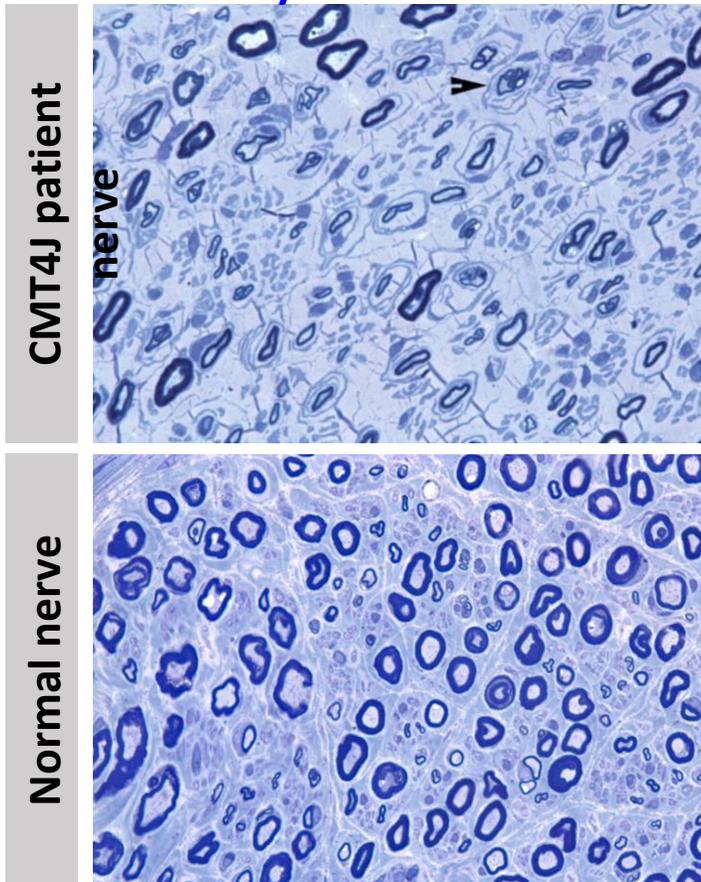
➤ Similar results with early treatment

Other CMT4: CMT4J

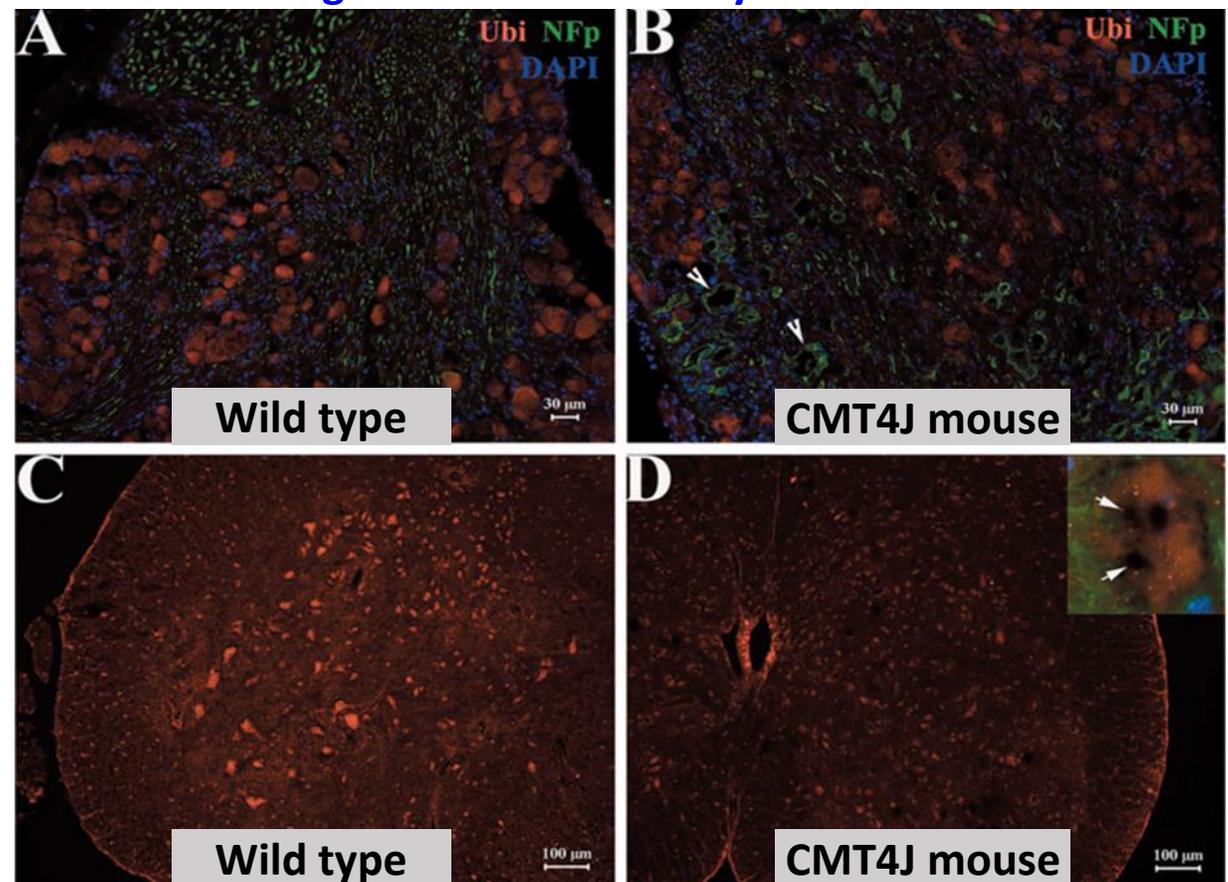
- CMT4J is the **0.3% of all CMT cases**
- Caused by **loss of function mutations in *FIG4* gene**
- *FIG4* is 5-phosphate serves in **endosome/lysosome pathway**

Zhang et al, 2008, Brain

Demyelination in CMT4J



Loss of large motor and sensory neurons in CMT4J



CMT4J gene therapy: AAV9-CBA/CMV-FIG4

Presa et al., 2021, J Clin Invest

Fig4-pale tremor (*plt*)

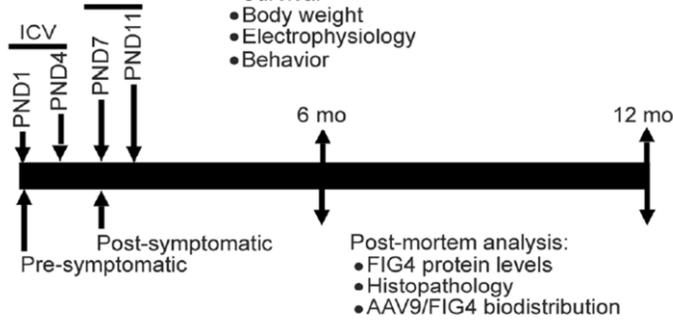


AAV9-CMV/CBA-FIG4 treatment:
postnatal day 1 or 4

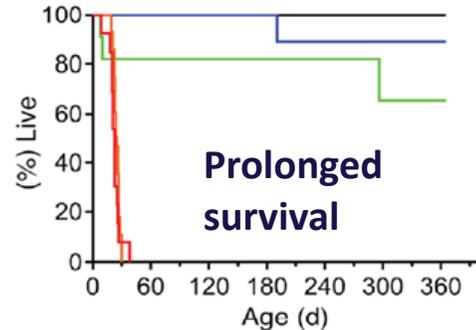
ICV: 5.4e11 vg/animal

Transposon insertion
in intron 18 of mu *Fig4*

In life monitoring:
• Survival
• Body weight
• Electrophysiology
• Behavior

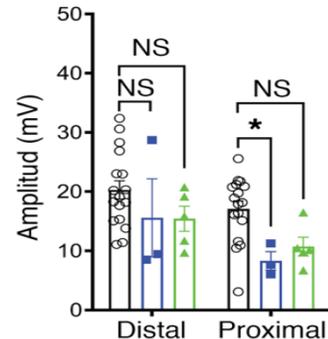


Post-mortem analysis:
• FIG4 protein levels
• Histopathology
• AAV9/FIG4 biodistribution

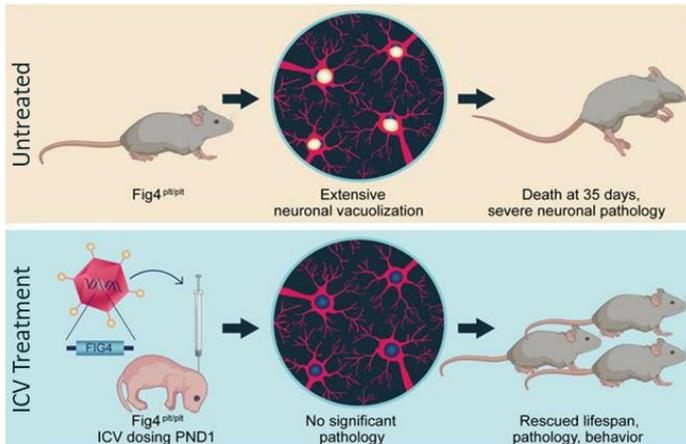


Prolonged survival

- *Fig4*^{+/+} untreated (n = 14)
- *Fig4*^{+/+} PND1 AAV9/FIG4 (n = 18)
- *Fig4*^{plt/plt} untreated (n = 13)
- *Fig4*^{plt/plt} vehicle (n = 10)
- *Fig4*^{plt/plt} PND1 AAV9/FIG4 (n = 12)
- *Fig4*^{plt/plt} PND4 AAV9/FIG4 (n = 11)



- *Fig4*^{+/+} untreated (n = 17)
- *Fig4*^{plt/plt} PND1 AAV9/FIG4 (n = 3)
- ▲ *Fig4*^{plt/plt} PND4 AAV9/FIG4 (n = 5)



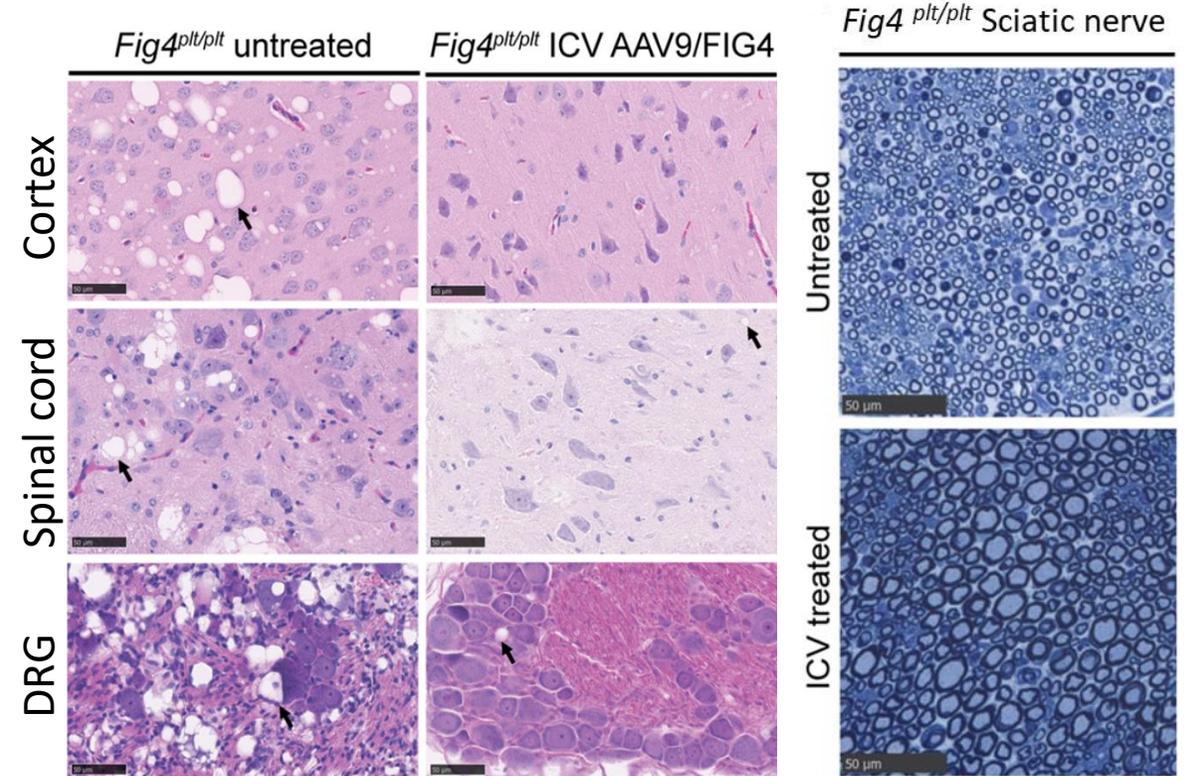
CMT4J gene therapy:

→ clinical trial, advocated by CureCMT4J, now part of Bespoke Gene Therapies Consortium

- P1 or P4 Tx → survival > 1 year, normal motor
- P7 or P11 Tx → prolonged survival and function **but incomplete**

Rescue of motor neurons

Improved myelination



Opportunities and challenges in gene therapy for neuropathies

Challenges in clinical translation

-  The **earlier the intervention** the more beneficial the treatment will be, but have **to get the dose right!**
-  CMT neuropathies **are non-life threatening diseases**, need to **ensure that “potential treatments” are safe**
-  **Need for sensitive clinical outcome measures** and **treatment responsive biomarkers**
-  Need **natural history data for ultra-rare forms, biomarkers** for all CMT types!
-  **Industry limitations**, financing clinical translation
-  **Regulatory requirements** and limitations

Opportunities

- ✓ **Strong advocacy Groups, Patient Associations**
- ✓ **Progress in Natural History Studies, biomarker discovery**
- ✓ **Collaborative Network** between scientists, clinicians, patients, industry
- ✓ **Increasing experience in clinical gene therapies and safety aspects**



- **Steven S. Scherer, MD, PhD, Univ. of Pennsylvania, PA**
- **Scott Harper, PhD, Nationwide Children's Columbus, OH**
- **Mary Reilly, MD and Alex Rossor, MD, Henrik Zetterberg, PhD, UCL, London**
- **Rita Horvath, MD, Univ. of Cambridge, UK**
- **John Svaren, PhD, Univ. of Wisconsin, Madison, WI**
- **Assumpció Bosch, PhD, Universitat Autònoma of Barcelona, Spain**
- **Steven Gray, PhD, UTSW, Dallas, TX**



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- **Melina Christou, PhD**
- **Styliani Theophanous, PhD**

Molecular Virology Department

- **Jan Richter, PhD**
- **Christina Tryphonos, PhD**
- **Christina Christodoulou, PhD**

